

Table 4

Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
6944	16822	27013	1.53	1.0E-06	AL1207878.1	EST_HUMAN	q123f06_x1 NCI_CGAP_Lym6 Homo sapiens cDNA clone IMAGE:1982435 3' similar to contains element MIR repetitive element;
7582	17433	27647	3.72	1.0E-06	UB25688.1	NT	Homo sapiens shox gene, alternatively spliced products, complete cds
7582	17433	27648	3.72	1.0E-06	UB25688.1	NT	Homo sapiens shox gene, alternatively spliced products, complete cds
7609	17460	27676	4.86	1.0E-06	AA132611.1	EST_HUMAN	z017e08_r1 Stratagene colon (#037204) Homo sapiens cDNA clone IMAGE:587174_5'
7637	17488		3.73	1.0E-06	AA449257.1	EST_HUMAN	z04d11_s1 Soares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:785493 3' similar to gb:D26129 RIBONUCLEASE PANCREATIC PRECURSOR (HUMAN);
7989	17839		1.52	1.0E-06	AL163203.2	NT	Homo sapiens chromosome 21 segment HS21C003
8912	18720		4.81	1.0E-06	AW890941.1	EST_HUMAN	RC4-NT0054-120500-012-b03 NT054-1 Homo sapiens cDNA
9440	19077	25280	2.99	1.0E-06	L78810.1	NT	Homo sapiens ADPATP carrier protein (ANT-2) gene, complete cds
9536	11844	21729	1.84	1.0E-06	AF184614.1	NT	Homo sapiens p47-phox (NCF1) gene, complete cds
9536	11844	21730	1.84	1.0E-06	AF184614.1	NT	Homo sapiens p47-phox (NCF1) gene, complete cds
357	10314	20134	1.49	9.0E-07	AF03529.1	NT	Homo sapiens glyican 3 (GP-C3) gene, partial cds and flanking repeat regions
357	10314	20135	1.49	9.0E-07	AF03529.1	NT	Homo sapiens glyican 3 (GP-C3) gene, partial cds and flanking repeat regions
8572	18440	28708	2.38	9.0E-07	AL163281.2	NT	Homo sapiens chromosome 21 segment HS21C081
4661	14547	24336	4.49	8.0E-07	AL120596.1	EST_HUMAN	q182g07_x1 Soares_NHMPu_S1 Homo sapiens cDNA clone IMAGE:1878876_3'
4661	14547	24337	4.49	8.0E-07	AL120596.1	EST_HUMAN	q182g07_x1 Soares_NHMPu_S1 Homo sapiens cDNA clone IMAGE:1878876_3'
5559	15504		7.45	8.0E-07	P21414	SWISSPROT	POL POLYPROTEIN [CONTAINS: PROTEASE ; REVERSE TRANSCRIPTASE ; ENDONUCLEASE]
6667	16547		7.89	8.0E-07	AF135416.1	NT	Homo sapiens UDP-glucuronosyltransferase gene, complete cds
8893	18703		7.2	8.0E-07	107770.1	EST_HUMAN	EST05660 Fetal brain, Stratagene (Cat#G35206) Homo sapiens cDNA clone HFBEN89
9055	18538		4.15	8.0E-07	AL163280.2	NT	Homo sapiens chromosome 21 segment HS21C080
1822	11719	21599	0.94	7.0E-07	AF167341.1	NT	Homo sapiens membrane interleukin 1 receptor accessory protein (IL1RAP) gene, exons 10 and 11
1870	11766	21641	2.36	6.0E-07	AW855558.1	EST_HUMAN	CM3-CT0277-221089-024-e11 CT0277 Homo sapiens cDNA
2442	12319	22217	2.41	6.0E-07	AF019413.1	NT	Homo sapiens HLA class III region containing tenascin X (tenascin-X) gene, partial cds; cytochrome P450 21-hydroxylase (CYP21B), complement component C4 (C4B) G11, helicase (SKI2W), RD, complement factor B (Bf), and complement component C2 (C2) genes, >
3891	13802		1.65	6.0E-07	P41479	SWISSPROT	HYPOTHETICAL 21.1 KD PROTEIN IN LEFL4-P23 INTERGENIC REGION
7279	17156	27351	1.35	6.0E-07	BF01867.1	EST_HUMAN	7g94f07_x1 NCI_CGAP_C016 Homo sapiens cDNA clone IMAGE:3314149_3' similar to TR:075920_075920_4F5L_;
9303	19053		1.81	6.0E-07	AW903222.1	EST_HUMAN	CM4-NN1029-250300-121-h12 NN1029 Homo sapiens cDNA
323	10284		3.55	5.0E-07	AB81893.1	EST_HUMAN	wh64f10_x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2386547_3'
1041	10569		2.9	5.0E-07	AA380630.1	EST_HUMAN	EST098615 Supt cells Homo sapiens cDNA 5' end
2995	12923		0.87	5.0E-07	AB831893.1	EST_HUMAN	wh64f10_x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2386547_3'

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
4547	14440	24223	1.44	5.0E-07	AF149774.1	NT	Homo sapiens NOD1 protein (NOD1) gene, exons 4 through 14 and complete cds
6203	15983	26096	1.71	5.0E-07	AI389981.1	EST_HUMAN	tg06b05.x1 NCI_CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2107953' similar to contains Alu repetitive element;contains element A3R repetitive element;
6203	15983	26097	1.71	5.0E-07	AI389981.1	EST_HUMAN	tg06b05.x1 NCI_CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2107953' similar to contains Alu repetitive element;contains element A3R repetitive element;
6348	16211	26373	15.93	5.0E-07	AV070885.1	EST_HUMAN	x31ef22.x1 NCI_CGAP_Br18 Homo sapiens cDNA clone IMAGE:2568362' similar to dbX15341 CYTOCHROME C OXIDASE POLYPEPTIDE VIA-LIVER (HUMAN);
7882	17782	28021	4.11	5.0E-07	AI908587.1	EST_HUMAN	CM-BT178-220498-014 BT178 Homo sapiens cDNA PRECURSOR
8805	18819	28909	4.31	5.0E-07	P11087	SWISSPROT	COLLAGEN ALPHA 1(I) CHAIN PRECURSOR
8863	18875			2.06	5.0E-07	AJ271735.1	NT
9712	18556			2.02	5.0E-07	AW382537.1	EST_HUMAN
3918	13827	23608	1.98	4.0E-07	AW009602.1	EST_HUMAN	QV0-CT0383-2/10400-204-b12 CT0383 Homo sapiens cDNA ws84n05.x1 NCI_CGAP_C03 Homo sapiens cDNA clone IMAGE:2504697'3'
7235	17112	27305	5.29	4.0E-07	AW419134.1	EST_HUMAN	xy49g11.x1 NCI_CGAP_Lj34.1 Homo sapiens cDNA clone IMAGE:2855548'3'
8308	18185	28432	3.84	4.0E-07	AI765528.1	EST_HUMAN	wi8hb08.x1 NCI_CGAP_Kid12 Homo sapiens cDNA clone IMAGE:2399703'3'
8308	18185	28433	3.84	4.0E-07	AI765528.1	EST_HUMAN	wi8hb08.x1 NCI_CGAP_Kid12 Homo sapiens cDNA clone IMAGE:2399703'3'
8552	18422			2.05	4.0E-07	BE001828.1	EST_HUMAN
434	10379	20200	5.01	3.0E-07	U19719.1	NT	Human microfibril-associated glycoprotein (MFAP2) gene, putative promoter region and alternatively spliced untranslated exons
568	10507	20314	2.38	3.0E-07	AJ271735.1	NT	Human microfibril-associated glycoprotein (MFAP2) gene, putative promoter region and alternatively spliced untranslated exons
1352	11258	21114	2.19	3.0E-07	M89149.1	NT	Human polymorphic microsatellite DNA
1608	11513			2.23	3.0E-07	M64857.1	NT
2000	11893			1.01	3.0E-07	AA526763.1	EST_HUMAN
2242	12126	22026	1.15	3.0E-07	M99149.1	NT	Human polymorphic microsatellite DNA
2420	12297	22194	17.25	3.0E-07	BE005077.1	EST_HUMAN	MRO-BN0115-020300-001-f11 BN0115 Homo sapiens cDNA PRECURSOR
2420	12297	22195	17.25	3.0E-07	BE005077.1	EST_HUMAN	MRO-BN0115-020300-001-f11 BN0115 Homo sapiens cDNA PRECURSOR
2998	12926	22118	0.85	3.0E-07	T84704.1	EST_HUMAN	yc50fr12.1 Soares fetal liver spleen /NFLS Homo sapiens cDNA clone IMAGE:111695'5'
3122	13047	22844	1.82	3.0E-07	P38739	SWISSPROT	HYPOTHETICAL 63.8 KD PROTEIN IN GUT1-RM1 INTERGENIC REGION PRECURSOR
4624	14512	24502	7.05	3.0E-07	AV660201.1	EST_HUMAN	AV660201 GLC Homo sapiens cDNA clone GI_CCCD01 3'
4654	14540	24329	0.89	3.0E-07	AI797236.1	EST_HUMAN	we86br12.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2347967'3'
4996	14871	24634	1.47	3.0E-07	T57850.1	EST_HUMAN	yc14h09.s1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:80705'3' similar to similar to gb:M62982 ARACHIDONATE 12-LIPOXYGENASE (HUMAN)
4996	14871	24635	1.47	3.0E-07	T57850.1	EST_HUMAN	yc14h09.s1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:80705'3' similar to similar to gb:M62982 ARACHIDONATE 12-LIPOXYGENASE (HUMAN)

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							PROTEIN-ARGININE DEIMINASE TYPE IV (PEPTIDYLARGININE DEIMINASE IV) (PAD-R4)
5476	15396	25462	10.26	3.0E-07	O88807	SWISSPROT	(PEPTIDYLARGININE DEIMINASE TYPE ALPHA)
6026	15930		5.32	3.0E-07	AAB15175.1	EST_HUMAN	cc04c10.s1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1339890 3'
6433	16294	26456	3	3.0E-07	AW797168.1	EST_HUMAN	QV1-LM0036-200300-115-g02 UM0036 Homo sapiens cDNA
8918	18726		3	3.0E-07	AF029308.1	NT	Homo sapiens chromosome 9 duplication of the T cell receptor beta locus and trypsinogen gene families
9939	19408		2.76	3.0E-07	AJ132352.1	NT	Rattus norvegicus mRNA for 45 kDa secretory protein, partial
26	10013	19807	2.62	2.0E-07	AF262988.1	NT	Homo sapiens TRF2 interacting telomeric RAP1 protein (RAP1) mRNA, complete cds
148	10122	19940	4.75	2.0E-07	L77569.1	NT	Homo sapiens DiGeorge syndrome critical region, telomeric end
148	10122	19941	4.75	2.0E-07	L77569.1	NT	Homo sapiens DiGeorge syndrome critical region, telomeric end
175	10146	19962	116.1	2.0E-07	U38849.1	NT	Fugu rubripes beta-cytoplasmic(vascular) actin gene, complete cds
731	10653	20495	1.71	2.0E-07	AF003530.1	NT	Homo sapiens homeobox protein CDX4 (CDX4) gene, complete cds and flanking repeat regions
731	10653	20496	1.71	2.0E-07	AF003530.1	NT	Homo sapiens homeobox protein CDX4 (CDX4) gene, complete cds and flanking repeat regions
923	10853	20701	3.11	2.0E-07	AA223260.1	EST_HUMAN	z108bb7_s1 Strategene NT2 neuronal precursor S37/230 Homo sapiens cDNA clone IMAGE:650869 3' similar to gb:31380 GLYCOPHORIN A PRECURSOR (HUMAN); contains Alu repetitive element;
929	10854	20702	3.38	2.0E-07	T63042.1	EST_HUMAN	yc15g04_s1 Strategene lung (#937210) Homo sapiens cDNA clone IMAGE:80790 3' similar to contains L1 repetitive element ;
1147	11060	20903	1.28	2.0E-07	Q25768	SWISSPROT	I/6 AUTOANTIGEN
1582	11486	21347	1.84	2.0E-07	Q05701	SWISSPROT	HYPOTHETICAL 72.5 KD PROTEIN C2F7.10 IN CHROMOSOME 1
3633	13547	23334	14.74	2.0E-07	AF25348.1	NT	Homo sapiens cavedilin 1 (CAV1) gene, exon 3 and partial cds
5078	14948	24723	0.97	2.0E-07	AW7070995.1	EST_HUMAN	xa05f07_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2567485 3' similar to WP:C38H2.1 CE00923 PROBABLE RABGAP DOMAINS ;
5078	14948	24724	0.97	2.0E-07	AW7070995.1	EST_HUMAN	xa05f07_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2567485 3' similar to WP:C38H2.1 CE00923 PROBABLE RABGAP DOMAINS ;
5198	15061	24826	0.95	2.0E-07	AL163301.2	NT	Homo sapiens chromosome 21 segment HS21/C101
5277	15199	24975	1.73	2.0E-07	AW898066.1	EST_HUMAN	RC3-NN00056-260400-021-g11 NN0066 Homo sapiens cDNA
6009	15914	26041	1.69	2.0E-07	A1208715.1	EST_HUMAN	qc95dd05_x1 Soares_festis_NHT Homo sapiens cDNA clone IMAGE:1889177 3'
6934	16812		3.91	2.0E-07	AV729390.1	EST_HUMAN	AV729390 HTC Homo sapiens cDNA clone HTCAE602 5'
7627	17478		1.61	2.0E-07	AL163303.2	NT	Homo sapiens chromosome 21 segment HS21/C103
7891	17741	27984	6.48	2.0E-07	AW892507.1	EST_HUMAN	CN4-NN0003-280300-124-e06 NN0003 Homo sapiens cDNA
9094	19557		1.48	2.0E-07	AI732462.1	EST_HUMAN	z185h11_x5 Strategene lung carcinoma 937218 Homo sapiens cDNA clone IMAGE:565029 3' similar to contains THR_b2 THR repetitive element ;
1086	11002		1.21	1.0E-07	AL163282.2	NT	Homo sapiens chromosome 21 segment HS21/C082

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2331	12212	22110	0.94	1.0E-07	P10263	SWISSPROT	RETROVIRUS-RELATED GAG POLYPYROTEIN (VERSION 1)
2403	12280	22177	0.94	1.0E-07	7549818	NT	Homo sapiens RAB, member of RAS oncogene family-like 2A (RABL2A), transcript variant 2, mRNA
2797	11410	21269	1.75	1.0E-07	P08256	SWISSPROT	GLYCOPROTEIN GPV
3684	11002		1.33	1.0E-07	AL163282.2	NT	Homo sapiens chromosome 21 segment HS21C082
4195	14095	23874	2.37	1.0E-07	AV71862	EST HUMAN	AV71862 GLC Homo sapiens cDNA clone GLCNIFO4 5'
4195	14095	23875	2.37	1.0E-07	AV71862	EST HUMAN	AV71862 GLC Homo sapiens cDNA clone GLCNIFO4 5'
6103	15997	26132	5.2	1.0E-07	BE047871.1	EST HUMAN	Iz43d06.y1 NCI_CGAP_Bm32 Homo sapiens cDNA clone IMAGE:2291339 5'
6103	15997	26133	5.2	1.0E-07	BE047871.1	EST HUMAN	Iz43d06.y1 NCI_CGAP_Bm32 Homo sapiens cDNA clone IMAGE:2291339 5'
6424	16235	26447	9.04	1.0E-07	N56081.1	EST HUMAN	yv43c07.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:2454843 3'
6782	16661	26850	2.81	1.0E-07	P97435	SWISSPROT	ENTEROPEPTIDASE (ENTEROKINASE)
6782	16661	26851	2.81	1.0E-07	P97435	SWISSPROT	ENTEROPEPTIDASE (ENTEROKINASE)
7180	17057	27246	3.24	1.0E-07	AA693576.1	EST HUMAN	Z51e10s1 Soares fetal liver spleen 1NFLS S1 Homo sapiens cDNA clone IMAGE:434346 3'
7714	17564	27790	2.37	1.0E-07	BF674524.1	EST HUMAN	602137714F1 NIH_MGC_83 Homo sapiens cDNA clone IMAGE:4274426 5'
7716	17566	27792	1.28	1.0E-07	AA386311.1	EST HUMAN	EST185054 Brain IV Homo sapiens cDNA
7980	17830		1.56	1.0E-07	AL163282.2	NT	Homo sapiens chromosome 21 segment HS21C082
9364	19533	25060	3.66	1.0E-07	BE048770.1	EST HUMAN	hr53ct11.x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:3132212 3' similar to TR:O95722 O95722
9497	19113		1.37	1.0E-07	X56467.1	NT	DJ1163J1.1;
9661	19222		4.61	1.0E-07	X55175.1	NT	H_sapiens ALAD gene for porphobilinogen synthase
7689	17539	27765	1.67	9.0E-08	AV734819.1	EST HUMAN	Human lambda-immunoglobulin constant region complex (germline)
8520	18392	28656	2.91	9.0E-08	AI891052.1	EST HUMAN	AV734819 cda Homo sapiens cDNA clone cdABFB06 5'
8922	18730	29025	3.91	9.0E-08	AL163301.2	NT	wn30a07.x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2446832 3' similar to contains OFR:t2
9316	19008		3.09	9.0E-08	A1251973.1	NT	OFR repetitive element;
591	12671		2.65	8.0E-08	AI911352.1	EST HUMAN	Homo sapiens partial sterin-1 gene
1034	10862		0.88	8.0E-08	BE795469.1	EST HUMAN	wd16b05.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:23228273 3'
3498	13415		1.43	8.0E-08	BE795469.1	EST HUMAN	601590133F1 NIH_MGC_7 Homo sapiens cDNA clone IMAGE:3943976 5'
7074	16951	27144	3.38	8.0E-08	AI7752367.1	EST HUMAN	cn15c02.x1 Normal Human Trabecular Bone Cells Homo sapiens cDNA clone NHTBC_cn15c02 tandem
7074	16951	27145	3.38	8.0E-08	AI7752367.1	EST HUMAN	cn15c02.x1 Normal Human Trabecular Bone Cells Homo sapiens cDNA clone NHTBC_cn15c02 tandem
7551	17402	27616	2.89	8.0E-08	AW970693.1	EST HUMAN	EST38278 MAGE resequences, MAGK Homo sapiens cDNA
8570	18438		2.39	8.0E-08	AF253417.1	NT	Hom sapiens microsomal epoxide hydrolase (EPHX1) gene, complete cds

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73	10057	19874	3.27	7.0E-08 Q022557	SWISSPROT	ANKYRIN 1 (ERYTHROCYTE ANKYRIN)	
1338	11244	21102	12.71	7.0E-08 X04809.1	NT	Rat mRNA for ribosomal protein L31	
3528	13444	23240	1.09	7.0E-08 P15305	SWISSPROT	DYNEIN HEAVY CHAIN (DYHC)	
3528	13444	23241	1.09	7.0E-08 P15305	SWISSPROT	DYNEIN HEAVY CHAIN (DYHC)	
8189	18075		4.63	7.0E-08 A1535743.1	EST_HUMAN	cong3.F11.A5 contorn Homo sapiens cDNA 3'	
8926	18734	29027	5.6	7.0E-08 U24070.1	NT	Rattus norvegicus Munc13-1 mRNA, complete cds	
9770	13444	23240	4.54	7.0E-08 P15305	SWISSPROT	DYNEIN HEAVY CHAIN (DYHC)	
9770	13444	23241	4.54	7.0E-08 P15305	SWISSPROT	DYNEIN HEAVY CHAIN (DYHC)	
9834	19334		1.89	7.0E-08 AJ31016.1	NT	Homo sapiens SCL gene locus	
800	10729	20569	2.84	6.0E-08 AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048	
800	10729	20570	2.84	6.0E-08 AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048	
2312	12193	22092	1.73	6.0E-08 BE144398.1	EST_HUMAN	MRO-HT0166-191198-004-g09 HT0166 Homo sapiens cDNA	
4155	14055	23829	0.99	6.0E-08 AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048	
8717	18534	28818	2.26	6.0E-08 P11369	SWISSPROT	RETRUVIUS-RELATED POL POLYPROTEIN [CONTAINS: REVERSE TRANSCRIPTASE ; ENDONUCLEASE]	
8819	18632		1.74	6.0E-08 AL163209.2	NT	Homo sapiens chromosome 21 segment HS21C009	
79	10063	19880	3.06	5.0E-08 AL163303.2	NT	Homo sapiens chromosome 21 segment HS21C03	
2189	12076	21981	1.97	5.0E-08 AA493851.1	EST_HUMAN	nh03b09.s1 NCI_CGAP_Thy1 Homo sapiens cDNA clone IMAGE:943193 similar to contains Alu repetitive element;	
5178	15042	24809	1.12	5.0E-08 Q06278	SWISSPROT	ALDEHYDE OXIDASE	
9057	18839		4.36	5.0E-08 P03681	SWISSPROT	COMPLEMENT C2 PRECURSOR (C3/C5 CONVERTASE)	
9245	18958	25317	1.94	5.0E-08 AW851878.1	EST_HUMAN	QY0-CT0225-131098t-034-a12 CT0225 Homo sapiens cDNA	
1724	11625	21493	1.1	4.0E-08 P25723	SWISSPROT	DORSAL-VENTRAL PATTERNING TOLLOID PROTEIN PRECURSOR	
1724	11625	21494	1.1	4.0E-08 P25723	SWISSPROT	DORSAL-VENTRAL PATTERNING TOLLOID PROTEIN PRECURSOR	
7278	17155	27350	1.41	4.0E-08 L42571.1	NT	Cricetulus griseus ribosomal transcription factor (UBF2) mRNA, complete cds	
7938	17788	28030	4.17	4.0E-08 A1050027.1	EST_HUMAN	an22d10.x1 Geesler Wilms tumor Homo sapiens cDNA clone IMAGE:1699411 3' similar to contains Alu repetitive element;contains element MER22 repetitive element;	
8274	18154		1.79	4.0E-08 AJ538617.1	NT	Homo sapiens mRNA for tRNA-associated antigenic protein (tRNA48 gene)	
8439	18313	28570	3.66	4.0E-08 BF592493.1	EST_HUMAN	602248024F1 NIH_MGC_62 Homo sapiens cDNA clone IMAGE:4335300 5'	
8439	18313	28571	3.66	4.0E-08 BF592493.1	EST_HUMAN	602248024F1 NIH_MGC_62 Homo sapiens cDNA clone IMAGE:4335300 5'	
9061	19680		1.54	4.0E-08 W76159.1	EST_HUMAN	zd65g03.1 Soares_fetal_heart_NbHH19W Homo sapiens cDNA clone IMAGE:345556 5' similar to contains L1.t1.L1 repetitive element;	
9703	19252		1.46	4.0E-08 A134353.1	EST_HUMAN	tb95af11.NC_CGAP_Co16 Homo sapiens cDNA clone IMAGE:2062076 3' similar to contains MER18.b3	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
5184	15048	24812	7.24	3.0E-08 AA191195.1	EST_HUMAN	zq45d05.r1 Stratagene hNT neuron #937/233 Homo sapiens cDNA clone IMAGE:632649 5'	
5184	15048	24813	7.24	3.0E-08 AA191195.1	EST_HUMAN	zq45d05.r1 Stratagene hNT neuron #937/233 Homo sapiens cDNA clone IMAGE:632649 5'	
5446	15367	25423	1.99	3.0E-08 BE018348.1	EST_HUMAN	bb79a10.y1 NIH_MGC_10 Homo sapiens cDNA clone IMAGE:3048570 5' similar to TR:Q32158 Q9Z158 SYNTAXIN 17.;	
6163	15120	24864	3.58	3.0E-08 AI792737.1	EST_HUMAN	qs16f11.y5 NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:1944045 5'	
6451	16312	26478	1.41	3.0E-08 AL163246.2	NT	Homo sapiens chromosome 21 segment HS21C046	
6558	16416		2.97	3.0E-08 AI436352.1	EST_HUMAN	lh93h10.x1 Soares_NSF_F8_9W_OT_PA_P_S11 Homo sapiens cDNA clone IMAGE:2126273 3' similar to TR:Q13537 Q13537 MER37 TRANSPOSABLE ELEMENT, COMPLETE CONSENSUS SEQUENCE.;	
9027	18819		11.6	3.0E-08 R18420.1	EST_HUMAN	yg02f04.r1 Soares_infant brain NIH3T3 Homo sapiens cDNA clone IMAGE:30948 5' similar to contains Alu repetitive element;	
199	10171		15.76	2.0E-08 AW302986.1	EST_HUMAN	xr87f06.x1 NCI_CGAP_Lu26 Homo sapiens cDNA clone IMAGE:2767139 3'	
223	10194		7.97	2.0E-08 AA425598.1	EST_HUMAN	2w48f07.11 Soares_total_fetus_Nb2hf8_9w Homo sapiens cDNA clone IMAGE:7733177 5' similar to contains NT	
488	10431	20245	1.48	2.0E-08 AF198349.1	EST_HUMAN	Alu repetitive element;contains element_MER15 repetitive element;	
644	10581	20396	10.93	2.0E-08 AW886438.1	EST_HUMAN	Gallus gallus Dach2 protein (Dach2) mRNA, complete cds	
644	10581	20397	10.93	2.0E-08 AW886438.1	EST_HUMAN	MRO-O10080-240200-001-908 OT10080 Homo sapiens cDNA	
974	10898		37.42	2.0E-08 BE280477.1	EST_HUMAN	MRO-O10080-240200-001-908 OT10080 Homo sapiens cDNA	
1319	11226	21082	1.93	2.0E-08 AL163247.2	NT	601153241F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:3138893 5'	
1707	11608		1.03	2.0E-08 BE734871.1	EST_HUMAN	Homo sapiens chromosome 21 segment HS21C047	
1811	11708		3.89	2.0E-08 AW270227.1	EST_HUMAN	601570463F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:3845199 5'	
2368	12248	22140	1.42	2.0E-08 AA731948.1	EST_HUMAN	xp43f11.x1 NCI_CGAP_HN11 Homo sapiens cDNA clone IMAGE:2743149 3'	
2446	12371		2.16	2.0E-08 KK0216.1	NT	nw6fh101.s1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1251409 3' similar to contains L1.3 L1 repetitive element;	
3171	13096	22901	6.02	2.0E-08 O12280	SWISSPROT	Sheep His-tRNA-GUG	
3171	13096	22902	6.02	2.0E-08 O12280	SWISSPROT	WNT-14 PROTEIN PRECURSOR	
3783	13695		1.66	2.0E-08 AW813620.1	EST_HUMAN	WNT-14 PROTEIN PRECURSOR	
3996	13893	23669	0.78	2.0E-08 U82668.1	NT	RC3-ST0197-161099-012403 ST0197 Homo sapiens cDNA	
4305	14203		3.35	2.0E-08 AA459040.1	EST_HUMAN	repetitive element; Homo sapiens shox gene, alternatively spliced products, complete cds	
4876	14756		2.89	2.0E-08 AW572881.1	EST_HUMAN	aa26c07.r1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:814380 5' similar to contains L1.2 L1 repetitive element;	
6722	16602	26791	1.31	2.0E-08 AA490121.1	EST_HUMAN	ab02g06.s1 Stratagene fetal retina S37202 Homo sapiens cDNA clone IMAGE:339674 3'	
8995	18799	26092	11.62	2.0E-08 BF58904.1	EST_HUMAN	naa32e09.x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:3257969 3'	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
1493	12686	21257	1.13	1.0E-08 P31792	SWISSPROT	POL POLYPROTEIN [CONTAINS: REVERSE TRANSCRIPTASE ; ENDONUCLEASE]	
2005	11897		2.92	1.0E-08 BE141959.1	EST_HUMAN	PM2-HT0130-150969-001-f12 HT0130 Homo sapiens cDNA	
3155	13080	22881	1.24	1.0E-08 BE246844.1	EST_HUMAN	TCBAP/D5232 Pediatric pre-B cell acute lymphoblastic leukemia Baylor-HGSC project=TCBAP5232 Homo sapiens cDNA clone TCBAP5232	
3155	13080	22882	1.24	1.0E-08 BE246844.1	EST_HUMAN	TCBAP/D5232 Pediatric pre-B cell acute lymphoblastic leukemia Baylor-HGSC project=TCBAP5232 Homo sapiens cDNA clone TCBAP5232	
5191	15054	24818	1.43	1.0E-08 AL163280.2	NT	Homo sapiens chromosome 21, segment HS21/C080	
5438	15358	25414	4.05	1.0E-08 AJ010770.1	NT	Homo sapiens hyperon gene, exons 1-50	
6975	16852	27045	1.84	1.0E-08 AJ015304.1	EST_HUMAN	035a05_s1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:1618736_3'	
8633	18498	28773	3.66	1.0E-08 AF044083.1	NT	Homo sapiens major histocompatibility locus class III region	
9434	19074		2.01	1.0E-08 X51755.1	NT	Human lambda-immunglobulin constant region complex (germline)	
9804	19316		6.3	1.0E-08 X51755.1	NT	Human lambda-immunglobulin constant region complex (germline)	
9894	19518		1.76	1.0E-08 BT75398.1	EST_HUMAN	MR4-ST0240-240700-013-g04 ST0240 Homo sapiens cDNA	
4149	14049	23823	2.98	9.0E-09 AL163279.2	NT	Homo sapiens chromosome 21, segment HS21/C079	
4149	14049	23824	2.98	9.0E-09 AL163279.2	NT	Homo sapiens chromosome 21, segment HS21/C079	
3444	13361		1.07	8.0E-09 BE012076.1	EST_HUMAN	RC5-BN1058-270400-031-C06 BN1058 Homo sapiens cDNA	
6307	16171	26328	6.12	8.0E-09 A1183500.1	EST_HUMAN	qd42e07_x1 Soares_fetal_heart_NbHH9W Homo sapiens cDNA clone IMAGE:1732164_3 similar to contains MSR1_t1 MSR1 repetitive element;	
6666	16546	26743	2.71	8.0E-09 AW900159.1	EST_HUMAN	CM0-NN1004-103000-273-e06 NN1004 Homo sapiens cDNA	
7196	17073		3.06	8.0E-09 AA938892.1	EST_HUMAN	op74d08_s1 SoaresNFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:1682575_3'	
3555	13469		1.68	7.0E-09 D86842.1	NT	Homo sapiens DNA for 3-ketoacyl-CoA thiolase beta-subunit of mitochondrial trifunctional protein, exon 2, 3	
4558	14450	24236	0.99	7.0E-09 D00649.1	NT	Homo sapiens gene for enteric smooth muscle gamma-actin, exon 5 end and flanking region	
7335	17203	27403	3.27	7.0E-09 L09709.1	NT	Human lysosomal membrane glycoprotein-2 (LAMP2) gene, 5' end and flanking region	
7850	17700	27945	1.84	7.0E-09 BE554850.1	EST_HUMAN	601111173Fe NIH_MGC_16 Homo sapiens cDNA clone IMAGE:335f1834_5'	
2107	11996		1.06	6.0E-09 AL0404039.1	EST_HUMAN	DKFZp434C0514_r1434 (synonym: ites3) Homo sapiens cDNA clone DKFZp434C0514_5'	
3941	13849	23625	1.06	6.0E-09 AA557940.1	EST_HUMAN	n117a11_s1 NCL_CGAP_HSC1 Homo sapiens cDNA clone IMAGE:1040924 similar to contains L1.t2 L1 repetitive element;	
4906	14786	24562	4.44	6.0E-09 BE169421.1	EST_HUMAN	PM1-HT0527-160200-001-h05 HT0527 Homo sapiens cDNA	
5305	15226	25030	8.19	6.0E-09 AW185784.1	EST_HUMAN	xn85h08_x1 SoaresNFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2701311_3'	
7294	17170	27370	2.26	6.0E-09 4503710	NT	Homo sapiens fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism) (FGFR3) mRNA	
7896	17746		4.06	6.0E-09 AF200923.2	NT	Homo sapiens testis-specific kinase substrate (TSKS) gene, complete cds	
1394	11299	21157	3.09	5.0E-09 BE149264.1	EST_HUMAN	RC2-HT0232-120200-014-h10 HT0232 Homo sapiens cDNA	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
5858	15764	25881	1.92	5.0E-09 AA359454.1	EST_HUMAN	EST68746 Fetal lung II Homo sapiens cDNA 5' end	
7811	17661	27901	2.69	5.0E-09 AW798667.1	EST_HUMAN	PM2-UN0053-240300-005-c09 UM0053 Homo sapiens cDNA	
510	10452		1.68	4.0E-09 AL163282.2	NT	Homo sapiens chromosome 21 segment HS21C082	
949	10873		2.31	4.0E-09 AL163285.2	NT	Homo sapiens chromosome 21 segment HS21C085	
1453	11368	21222	0.95	4.0E-09 9558718 NT	EST_HUMAN	Homo sapiens hypothetical protein (AF038169), mRNA	
2370	12269	22151	6.36	4.0E-09 AA350878.1	EST_HUMAN	EST8385 Infant brain Homo sapiens cDNA 5' end similar to heat shock protein, 30 kDa	
2303	12184	22082	3.82	3.0E-09 BE222239.1	EST_HUMAN	hu09e05_X1 NC1_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3166120 3' similar to contains MER18.13	
2506	12380	22270	1.25	3.0E-09 BE222239.1	EST_HUMAN	MER18 repetitive element;	
2614	12462	22371	1.1	3.0E-09 P23249	SWISSPROT	PROTEIN MOV-10	
3287	13208	23008	1.1	3.0E-09 BE222239.1	EST_HUMAN	hu09e05_X1 NC1_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3166120 3' similar to contains MER18.13	
4329	14226	24008	3.22	3.0E-09 AF76325.1	NT	Homo sapiens eukaryotic initiation factor 4A1 (EIF4A1) gene, partial cds	
4411	14505	24088	1.54	3.0E-09 Q9J3R5	SWISSPROT	258.1 KDA PROTEIN C21ORF5 (KIAA0933)	
7884	17734	27978	1.73	3.0E-09 AL163247.2	NT	Homo sapiens chromosome 21 segment HS21C047	
8384	18261	28510	3.8	3.0E-09 BF1_09943.1	EST_HUMAN	7172c08_x1 Scarecs_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE:3527030 3'	
8384	18261	28511	3.8	3.0E-09 BF1_09943.1	EST_HUMAN	7172c08_x1 Scarecs_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE:3527030 3'	
795	10724		0.93	2.0E-09 X1674.1	NT	H.sapiens PADPRP-1 gene for NAD(+) ADP-ribosyltransferase	
1237	11144	20995	5.23	2.0E-09 AL163284.2	NT	Homo sapiens chromosome 21 segment HS21C084	
1637	11541		10.52	2.0E-09 AL118573.1	EST_HUMAN	DKFZp761B1710_r1_761 (synonym: hamy2) Homo sapiens cDNA clone DKFZp761B1710 5'	
2278	12162	22059	2.25	2.0E-09 Q9J3R5	SWISSPROT	258.1 KDA PROTEIN C21ORF5 (KIAA0933)	
3888	13769	23561	3.65	2.0E-09 O60241	SWISSPROT	BRAIN-SPECIFIC ANGIOGENESIS INHIBITOR 2 PRECURSOR	
6405	16266	26428	8.9	2.0E-09 AA4461430.1	EST_HUMAN	zg63hd6_r1 Scarecs_fetal_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:796187 5' similar to contains Alu repetitive element;	
7055	16932	27122	1.37	2.0E-09 A1271735.1	NT	Homo sapiens Xq pseudoautosomal region; segment 1/2	
8569	18457	28726	2.11	2.0E-09 AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048	
9576	10724		11.53	2.0E-09 X1674.1	NT	H.sapiens PADPRP-1 gene for NAD(+) ADP-ribosyltransferase	
9640	19749		1.62	2.0E-09 AA226070.1	EST_HUMAN	nc11c02_r1 NC1_CGAP_P11 Homo sapiens cDNA clone IMAGE:1007810 similar to contains Alu repetitive element;	
1093	11009	20850	2.48	1.0E-09 5031624 NT		Homo sapiens CCAAT-T-box-binding transcription factor (CBF2) mRNA	
1093	11009	20851	2.48	1.0E-09 5031624 NT		Homo sapiens CCAAT-T-box-binding transcription factor (CBF2) mRNA	
1616	11520		0.95	1.0E-09 A1229041.1	NT	Homo sapiens 969 kb contig between AML1 and CBR1 on chromosome 21q22, segment 1/3	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal Value	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
2453	12330		0.94	1.0E-09 A 356086.1	EST_HUMAN	qj64e11.X1 NCI_CGAP_Bm25 Homo sapiens cDNA clone IMAGE:2068112 3' similar to contains MER12.t2	MER12 repetitive element;
2860	12788	22580	1.51	1.0E-09 U 80017.1	NT	Homo sapiens basic transcription factor 2 p44 (bit2p44) gene, partial cds, neuronal apoptosis inhibitory protein (napip) and survival motor neuron protein (smn) genes, complete cds	
2895	12822	22614	3.25	1.0E-09 M 28699.1	NT	Homo sapiens nucleolar phosphoprotein B23 (NPM1) mRNA, complete cds	
2895	12822	22615	3.25	1.0E-09 M 28699.1	NT	Homo sapiens nucleolar phosphoprotein B23 (NPM1) mRNA, complete cds	
2949	12876	22674	1.23	1.0E-09 P 11799	SWISSPROT	MYOSIN LIGHT CHAIN KINASE, SMOOTH MUSCLE AND NON-MUSCLE ISOZYMES (MLCK) [CONTAINS: TELOKIN]	
3002	12930	22722	0.78	1.0E-09 BE 355440.1	EST_HUMAN	60 05802F1 NIH_MGC_10 Homo sapiens cDNA clone IMAGE:3445177 5'	
4692	14578		4.26	1.0E-09 AA 719297.1	EST_HUMAN	zh85b03.s1 Soares_Pheal_gland_N3HPG Homo sapiens cDNA clone IMAGE:414029 3' similar to contains Alu repetitive element; contains element MER22 repetitive element;	
5560	15476	25549	1.37	1.0E-09 U 07000.1	NT	Human breakpoint cluster region (BCR) gene, complete cds	
5736	15644	25749	3.21	1.0E-09 P 28694	SWISSPROT	CIRCUITSPOROZOITE PROTEIN PRECURSOR (CS)	
7911	17761		3.1	1.0E-09 AL 163283.2	NT	Homo sapiens chromosome 21 segment HS21C083	
9478	19659	24996	2.14	1.0E-09 114 18127	NT	Homo sapiens GTP binding protein 1 (GTPBP1), mRNA	
1287	11195	21043	1.8	9.0E-10 AW 867740.1	EST_HUMAN	MR0-SN0040-050500-002-c07 SNO040 Homo sapiens cDNA	
2803	12733	22532	6.03	9.0E-10 A 870071.1	EST_HUMAN	we78h03.x1 Soares_Dieckgraefe_colon_NHCD Homo sapiens cDNA clone IMAGE:2347633 3' similar to SW:RI 29_HUMAN P4 9114 60S RIBOSOMAL PROTEIN L29, contains element P R5 repetitive element;	
6087	16032	26172	4.63	9.0E-10 A 452982.1	EST_HUMAN	jj46b09.x1 Soares_NSF_F8_9W_OT_PA_P_S1Homo sapiens cDNA clone IMAGE:2144537 3' similar to TR:000312_000312_PUTATIVE_P150.;	
141	10115	19935	9.43	8.0E-10 U 63630.2	NT	Homo sapiens MCM4 (MCM4) and DNA-PKcs (PRKDC) genes, partial cds	
3300	13222	23022	0.87	8.0E-10 BE 80748.1	EST_HUMAN	QV1-BT0631-150200-071-01 BT0631 Homo sapiens cDNA	
4106	14006	23782	2.82	8.0E-10 AA 376852.1	EST_HUMAN	EST89564 Small intestine Homo sapiens cDNA 5' end	
7725	17575		2.32	8.0E-10 U 36308.2	NT	Homo sapiens chromosome 21 segment HS21C080	
9005	18808	29100	2.31	8.0E-10 AL 163280.2	NT	Homo sapiens lens major intrinsic protein (MIP) gene, complete cds	
686	10619	20444	12.45	7.0E-10	7706225 NT	Homo sapiens TPA inducible protein (LOC55586), mRNA	
686	10619	20445	12.45	7.0E-10	7706225 NT	Homo sapiens TPA inducible protein (LOC55586), mRNA	
1605	11510	21371	1.87	7.0E-10 Q 13342	SWISSPROT	LYSP100 PROTEIN (LYMPHOID-RESTRICTED HOMOLOG OF SP100)	
1974	11867		1.01	7.0E-10 P 08548	SWISSPROT	LINE-1 REVERSE TRANSCRIPTASE HOMOLOG	
2512	12386		16.88	7.0E-10 P 08547	SWISSPROT	LINE-1 REVERSE TRANSCRIPTASE HOMOLOG	
3049	12976	22768	2.65	7.0E-10 X 00856.1	NT	H_sapiens DHFR gene, exon 3	
5754	15692	25769	3.98	7.0E-10 AA 345220.1	EST_HUMAN	EST51247 Gall bladder II Homo sapiens cDNA 5' end	
6514	16373		1.39	7.0E-10 P 35084	SWISSPROT	DNA-DIRECTED RNA POLYMERASE II LARGEST SUBUNIT	

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895	10821	20666	2.81	6.0E-10 AJ400877.1	NT	Homo sapiens ASCL3 gene, CEGP1 gene, C11orf14 gene, C11orf15 gene, C11orf17 gene
2644	12511	22402	1.47	6.0E-10 AJ424405.1	EST_HUMAN	It02407_x1 NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:2096021 3'
4634	14522		2.51	6.0E-10 AW853719.1	EST_HUMAN	RC3-CT0254-031086-012-g12 CT0254 Homo sapiens cDNA
9092	18865		1.79	6.0E-10 AW971923.1	EST_HUMAN	EST38-012 MAGE resequences, MAGL Homo sapiens cDNA RC3-NN0070-110800-014-h07 NN0070 Homo sapiens cDNA
9947	19597		3.54	6.0E-10 BE599410.1	EST_HUMAN	DKFZp434N219_t1 434 (synonym: hies3) Homo sapiens cDNA clone DKFZp434N219 5'
744	10675		4.5	5.0E-10 AL046804.1	EST_HUMAN	SWISSPROT HYPOTHETICAL PROTEIN
3430	13347	23152	1.48	5.0E-10 Q01033	SWISSPROT	6016222-184F1 NIH_MGC_75 Homo sapiens cDNA clone IMAGE:4042413 5'
6334	16197		1.82	5.0E-10 BF105159.1	EST_HUMAN	7496 27570 1.79 P34678 SWISSPROT HYPOTHETICAL 67.9 KD PROTEIN ZK68.8 IN CHROMOSOME III
7496	17366	27570	1.79	5.0E-10 P34678	SWISSPROT	7496 27571 1.79 P34678 SWISSPROT HYPOTHETICAL 67.9 KD PROTEIN ZK68.8 IN CHROMOSOME III
106	10087		0.99	4.0E-10 AL221083.1	EST_HUMAN	cg03f05_x1 Soares, placenta, 86weeks. 2NbHP8t0gVN Homo sapiens cDNA clone IMAGE:1759049 3' similar to contains LTR8_b2 LTR8 repetitive element;
1951	11846	21732	1.4	4.0E-10 AW594709.1	EST_HUMAN	hg55g03_x1 NCI_CGAP_GC8 Homo sapiens cDNA clone IMAGE:2949844 3' similar to contains Alu repetitive element;
2527	12401	22292	6.09	4.0E-10 AL163303.2	NT	Hom sapiens chromosome 21 segment HS21C103
6259	16125	26278	19.23	4.0E-10 AF224669.1	NT	Hom sapiens mannosidase, beta A, lysosomal (MANBA) gene, and ubiquitin-conjugating enzyme E2D 3 (UBE2D3) genes, complete cds
8436	18310		7.7	4.0E-10 AF003528.1	NT	Hom sapiens X-linked anhidrotic ectodermal dysplasia protein gene (EDA), exon 2 and flanking repeat regions
897	10822	20668	1.72	3.0E-10 N36113.1	EST_HUMAN	yg32t06_s1 Soares melanocyte 2NbHM Homo sapiens cDNA clone IMAGE:272963 3' similar to contains L1_tf1L1 repetitive element;
1329	11236		6.63	3.0E-10 AY005150.1	NT	Hom sapiens extracellular glycoprotein lactitin precursor, gene, complete cds
4435	14330	24117	1.1	3.0E-10 AL163203.2	NT	Hom sapiens chromosome 21 segment HS21C003
4435	14330	24118	1.1	3.0E-10 AL163203.2	NT	Hom sapiens chromosome 21 segment HS21C003
5765	15672	25779	2.83	3.0E-10 P20350	SWISSPROT RHOMBOID PROTEIN (VENNET PROTEIN)	ba76d08_v1 NIH_MGC_20 Homo sapiens cDNA clone IMAGE:2906319 5'
5833	15739	25851	3.27	3.0E-10 BE302970.1	EST_HUMAN	AV743302 CB Homo sapiens cDNA clone CBFBGD08 5'
6562	16420	26599	1.31	3.0E-10 AV743302.1	EST_HUMAN	6562 16420 26600 1.31 3.0E-10 AV743302.1 EST_HUMAN AV743302 CB Homo sapiens cDNA clone CBFBGD08 5'
7067	16944	27136	1.58	3.0E-10 H87208.1	EST_HUMAN	ys74b12_s1 Soares retina N2b4HR Homo sapiens cDNA clone IMAGE:220511 3' similar to contains MER29 repetitive element;
7233	17110	27302	1.47	3.0E-10 AW850731.1	EST_HUMAN	II_3-CT0219-160200-064-B06 CT0219 Homo sapiens cDNA
7233	17110	27303	1.47	3.0E-10 AW850731.1	EST_HUMAN	II_3-CT0219-160200-064-B06 CT0219 Homo sapiens cDNA
7979	17829		2.56	3.0E-10 T68891.1	EST_HUMAN	ycf1ef12_r1 Stratagene lung (#37210) Homo sapiens cDNA clone IMAGE:80398 5'

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Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
8026	17876			1.54	3.0E-10 A/1769294.1	EST_HUMAN	nc23g03.s1 NC1_C3'AP_GCB1 Homo sapiens cDNA clone IMAGE:1289908_3'
9737	19271	25227		2.03	3.0E-10 BE179517.1	EST_HUMAN	IL3-HT0618-110500-136-E07 HT0618 Homo sapiens cDNA
32	10019	19814		1.43	2.0E-10 P48988	SWISSPROT	MAJOR CENTROMERE AUTOANTIGEN B (CENTROMERE PROTEIN B) (CENP-B)
32	10019	19815		1.43	2.0E-10 P48988	SWISSPROT	MAJOR CENTROMERE AUTOANTIGEN B (CENTROMERE PROTEIN B) (CENP-B)
1855	11751			5.91	2.0E-10 U80017.1	NT	Homo sapiens basic transcription factor 2 p44 (hif2p44) gene, partial cds, neuronal apoptosis inhibitory protein (napip) and survival motor neuron protein (smn) genes, complete cds
5547	15463			2.41	2.0E-10 Q28640	SWISSPROT	(HPRG)
							Homo sapiens cytochrome P450 polypeptide 43 (CYP3A43) gene, partial cds; cytochrome P450 polypeptide 4 (CYP3A4) and cytochrome P450 polypeptide 7 (CYP3A7) genes, complete cds; and cytochrome P450 polypeptide 5 (CYP3A5) gene, partial cds
5787	15693	25801		1.71	2.0E-10 AF280107.1	NT	
6367	16230	26389		5.79	2.0E-10 BE791082.1	EST_HUMAN	601588208F1 NIH MCG_7 Homo sapiens cDNA clone IMAGE:3940824_5'
1492	11397			3.56	1.0E-10 AW867767.1	EST_HUMAN	MRO-SN0038-290300-001-f01 SN0038 Homo sapiens cDNA
1689	11493	21353		3.14	1.0E-10 AV652123.1	EST_HUMAN	AV652123 GLC Homo sapiens cDNA clone GLC CXA11_3'
2537	12411			2.4	1.0E-10 AW852001.1	EST_HUMAN	QV0-C0225-191198-058-e08 C70225 Homo sapiens cDNA
3456	13372	23178		0.89	1.0E-10 AW832812.1	EST_HUMAN	QV2-TT0003-161199-013-g10 TT0003 Homo sapiens cDNA
3770	13411			0.91	1.0E-10 AL041685.1	EST_HUMAN	DKFZp434N1317_r1434 (synonym: htes2) Homo sapiens cDNA clone DKFZp434N1317_5'
3933	13842			5.44	1.0E-10 AF213884.1	NT	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (NFkB1) gene, complete cds
4036	13939	23716		4.51	1.0E-10 U52111.2	NT	Homo sapiens X28 region near ALD locus containing dual specificity phosphatase 9 (DUSP9), ribosomal protein L18a (RPL18a), Ca2+/Calmodulin-dependent protein kinase I (CAMK1), creatine transporter (CRTTR), CDM protein (CDM), adrenoleukodystrophy protein >
4036	13939	23717		4.51	1.0E-10 U52111.2	NT	Homo sapiens X28 region near ALD locus containing dual specificity phosphatase 9 (DUSP9), ribosomal protein L18a (RPL18a), Ca2+/Calmodulin-dependent protein kinase I (CAMK1), creatine transporter (CRTTR), CDM protein (CDM), adrenoleukodystrophy protein >
4042	13945	23724		2.25	1.0E-10 AB031069.1	NT	Homo sapiens PCCX1 mRNA for protein containing CXXC domain 1, complete cds
4079	13981			2.06	1.0E-10 M30629.1	NT	Human pregnancy-specific glycoprotein beta-1 (SP-1) mRNA, last exon H_sapiens DMA, DMB, HLA-Z1, IPP2, LMP2, TAP1, LMP7, TAP2, DOB, DQB2 and RING3, 9, 13 and 14 genes
5145	15012			0.93	1.0E-10 X87344.1	NT	
7859	17709			4.59	1.0E-10 AA081868.1	EST_HUMAN	ZN23g06_r1 Stratagene neuroepithelium NT2RAM/11937234 Homo sapiens cDNA clone IMAGE:548314_5'
8288	18165	28408		3.4	1.0E-10 AI038280.1	EST_HUMAN	085f03_x1 Scores fetal liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:1672661_3'
9037	15012			1.43	1.0E-10 X87344.1	NT	H_sapiens DMA, DMB, HLA-Z1, IPP2, LMP2, TAP1, LMP7, TAP2, DOB, DQB2 and RING3, 9, 13 and 14 genes
9084	18859			1.3	1.0E-10 AA397885.1	EST_HUMAN	Zz86b10_r1 Scores testis_NHT Homo sapiens cDNA clone IMAGE:729271_5'

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261	10226	20042	1.2	9.0E-11 BE145600.1	EST_HUMAN	IL2-HT0203-291099-016-c08 HT0203 Homo sapiens cDNA clone DKFZp547D225_11 5'47 (synonym: hfbr1) Homo sapiens cDNA clone DKFZp547D225 5'		
2057	11947	21843	5.43	9.0E-11 AL134395.1	EST_HUMAN	DKFZp547D225_11 5'47 (synonym: hfbr1) Homo sapiens cDNA clone DKFZp547D225 5'		
2057	11947	21844	5.43	9.0E-11 AL134395.1	EST_HUMAN	DKFZp547D225_11 5'47 (synonym: hfbr1) Homo sapiens cDNA clone DKFZp547D225 5'		
3337	13257	23064	—	2.25	9.0E-11 AL134395.1	EST_HUMAN	DKFZp547D225_11 5'47 (synonym: hfbr1) Homo sapiens cDNA clone DKFZp547D225 5'	
3337	13257	23065	2.25	9.0E-11 AL134395.1	EST_HUMAN	DKFZp547D225_11 5'47 (synonym: hfbr1) Homo sapiens cDNA clone DKFZp547D225 5'		
4400	14295	24079	1.09	9.0E-11 AA1775985.1	EST_HUMAN	aer78f01_s1 Strategene schizo brain S11 Homo sapiens cDNA clone IMAGE:970287 3'		
5421	15342	—	4	9.0E-11 BE779780.1	EST_HUMAN	RC6-BT0627-140200-011-E06 BT0627 Homo sapiens cDNA		
9410	19052	25312	2.69	9.0E-11 C16635.1	EST_HUMAN	C16635 Clontech human aorta polyA+ mRNA (#6572) Homo sapiens cDNA clone GEN-5066B008 5'		
3078	13005	—	9.97	8.0E-11 H19971.1	EST_HUMAN	yn53f11_s1 Soares adult brain N265HB557 Homo sapiens cDNA clone IMAGE:172173 3' similar to contains L1 repetitive element;		
3881	13792	23580	0.83	8.0E-11 A1478617.1	EST_HUMAN	lrm54c05_x1 NCI CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2161936 3'		
3957	13885	23641	4.03	8.0E-11 IN23712.1	EST_HUMAN	yw46e06_s1 Weizmann Olfactory Epithelium Homo sapiens cDNA clone IMAGE:255298 3'		
1431	11336	21202	2.11	7.0E-11 AA530642.1	EST_HUMAN	EST34392 Embryo, 6 week Homo sapiens cDNA 5' end		
6949	16827	27020	2.55	7.0E-11 AF_63584.1	NT	Homo sapiens SNCA isoform (SNCA) gene, complete cds, alternatively spliced		
7873	17723	—	1.22	7.0E-11 P11369	SWISSPROT	RETRORVIRUS-RELATED POL POLYPROTEIN [CONTAINS: REVERSE TRANSCRIPTASE ; ENDONUCLEASE]		
9545	19145	—	1.31	7.0E-11 AV701656.1	EST_HUMAN	AV701656 ADB Homo sapiens cDNA clone ADBABC09 5'		
406	10352	20180	5.16	6.0E-11 M53270.1	NT	Human matrix Glu protein (MGP) gene, complete cds		
406	10352	20181	5.16	6.0E-11 M53270.1	NT	Human matrix Glu protein (MGP) gene, complete cds		
6529	16388	26563	3.58	6.0E-11 P08547	SWISSPROT	LINE-1 REVERSE TRANSCRIPTASE HOMOLOG		
6871	16750	26945	6.49	6.0E-11 AV727859.1	EST_HUMAN	AV727859 HTC Homo sapiens cDNA clone HTCAASC06 5'		
11	9997	19788	0.92	5.0E-11 AL163283.2	NT	Homo sapiens chromosome 21 segment HS21C083		
3320	9997	19788	1.48	5.0E-11 AL163283.2	NT	Homo sapiens chromosome 21 segment HS21C083		
4135	14035	23811	1.49	5.0E-11 P48034	SWISSPROT	ALDEH'DE OXIDASE		
5926	15831	25954	1.69	5.0E-11 AL163213.2	NT	Homo sapiens chromosome 21 segment HS21C013		
6446	16307	26472	14.59	5.0E-11 114116799	NT	Homo sapiens protocadherin beta 3 (PCDHB3), mRNA		
8975	18780	29072	1.79	5.0E-11 A1289880.1	NT	Homo sapiens KIAA0551 gene (partial), XT3 gene and LZTF1 gene		
1380	11285	—	1.4	4.0E-11 AA436042.1	EST_HUMAN	zq1b12_f1 Soares testis NHT Homo sapiens cDNA clone IMAGE:730559 5'		
2760	12622	22514	8.45	4.0E-11 BE885900.1	EST_HUMAN	601507331FH NIH_MGC_71 Homo sapiens cDNA clone IMAGE:3909295 5'		
4518	14411	24196	1.37	4.0E-11 D44666.1	EST_HUMAN	HUMSUPY069 Human brain cDNA Homo sapiens cDNA clone 069		
5897	15803	25927	2.94	4.0E-11 P20985	SWISSPROT	PRE-mRNA SPLICING FACTOR RNA HELICASE PRP2		
6362	16225	—	3.66	4.0E-11 AF224693.1	NT	Homo sapiens mannosidase, beta A, lysosomal (MANBA) gene, and ubiquitin-conjugating enzyme E2D 3 (UBE2D3) genes, complete cds		
7405	17272	—	1.59	4.0E-11 BE149425.1	EST_HUMAN	RC1-HT0286-210100-013-108 HT0286 Homo sapiens cDNA		

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9611	19190	25252	1.93	4.0E-11	11545732	NT	Homo sapiens SH3 domain binding protein 1 (SH3BP1), mRNA
1476	11381	21245	2.15	3.0E-11	6679077	NT	Mus musculus expressed in non-metastatic cells 2, protein (NM_023B) (Nm2e2), mRNA
4179	14079		1.45	3.0E-11	AA309248.1	EST_HUMAN	EST180120 Liver, hepatocellular carcinoma Homo sapiens cDNA 5' end
944	10869	20716	1.58	2.0E-11	A1150502.1	EST_HUMAN	qf36c04_x1 Soares_tests_NHT Homo sapiens cDNA clone IMAGE:1752102 3' similar to contains MER10.13 MER10 repetitive element;
1168	11080	20926	3.64	2.0E-11	R24807.1	EST_HUMAN	ygj3er12.1 Soares_infant brain 1N13 Homo sapiens cDNA clone IMAGE:351144 5'
1168	11080	20927	3.64	2.0E-11	R24807.1	EST_HUMAN	ygj3er12.1 Soares_infant brain 1N13 Homo sapiens cDNA clone IMAGE:351144 5'
1596	11560	21359	3.91	2.0E-11	L17432.1	NT	Gallus gallus rho-globin, beta-H globin, epsilon-globin, epsilon-globin, and olfactory receptor-like protein COR3'beta (COR3'beta) genes, complete cds
1596	11560	21360	3.91	2.0E-11	L17432.1	NT	Gallus gallus rho-globin, beta-H globin, beta-A globin, epsilon-globin, and olfactory receptor-like protein COR3'beta (COR3'beta) genes, complete cds
1600	11565	21365	1.04	2.0E-11	A1126371.1	EST_HUMAN	qcs1c10_x1 Soares_pregnant uterus NbHPU Homo sapiens cDNA clone IMAGE:1713138 3' similar to gbjL02932 PEROXISOME PROLIFERATOR ACTIVATED RECEPTOR ALPHA (HUMAN);contains L1:t1 L1 repetitive element;
2737	12599	22493	1.11	2.0E-11	AF087913.1	NT	Human endogenous retrovirus HERV-P-T47D
3160	13085	22889	4.23	2.0E-11	P10263	SWISSPROT	RETROVIRUS-RELATED GAG POLYPROTEIN (VERSION 1)
3284	13205	23005	0.87	2.0E-11	A1478617.1	EST_HUMAN	Im54c02_x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2161936 3'
3452	13368		0.95	2.0E-11	AF020503.1	NT	Homo sapiens FRA3B common fragile region, diadenosine triphosphate hydrolase (FHIT) gene, exon 5
4501	14395		0.97	2.0E-11	A1163227.2	NT	Homo sapiens chromosome 21 segment HS21C027
4851	14732		5.46	2.0E-11	BE062558.1	EST_HUMAN	QV2-BT0258-261098-014-#01 BT0258 Homo sapiens cDNA EST178226 Colon carcinoma (HCC) cell line Homo sapiens cDNA 5' end similar to alpha-2-
4961	14836	24604	1.32	2.0E-11	AA307331.1	EST_HUMAN	macroglobulin nc83h05_r1 NCI CGAP_GC1 Homo sapiens cDNA clone IMAGE:797435 5' similar to SWPR16_YEAST
5810	15715	25828	1.83	2.0E-11	AA581028.1	EST_HUMAN	P15938 PRE-mRNA SPlicing FACTOR RNA HELICASE PRP16. ;
7318	17194		1.43	2.0E-11	AF026308.1	NT	Homo sapiens chromosome 9 duplication of the T cell receptor beta locus and trypsinogen gene families OLFACTORY RECEPTOR 51(Olfactory RECEPTOR-LIKE PROTEIN OLF1)
7900	17750	27989	4.61	2.0E-11	Q13606	SWISSPROT	zK27g02_s1 Soares_pregnant uterus NbHPU Homo sapiens cDNA clone IMAGE:471794 3'
8457	18330	28591	2.07	2.0E-11	AA035369.1	EST_HUMAN	zK27g02_s1 Soares_pregnant uterus NbHPU Homo sapiens cDNA clone IMAGE:471794 3'
8457	18330	28592	2.07	2.0E-11	AA035369.1	EST_HUMAN	zK27g02_s1 Soares_pregnant uterus NbHPU Homo sapiens cDNA clone IMAGE:471794 3'
9160	19679		1.29	2.0E-11	AA704495.1	EST_HUMAN	zJ77e03_s1 Soares_fetal liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:460924 3'
9192	18927		2.25	2.0E-11	AV842143.1	EST_HUMAN	RC0-CN0027_210100-011-c01 CN0027 Homo sapiens cDNA
9216	18943	25356	1.98	2.0E-11	BF577859.1	EST_HUMAN	CN2-TN0140-070900-372-g01 TN0140 Homo sapiens cDNA
9477	19103		1.89	2.0E-11	D28217.2	NT	Homo sapiens mRNA for KIAA0027 protein, partial cds

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9627	19201		2.38	2.0E-11	P08547	SWISSPROT	LINE-1 REVERSE TRANSCRIPTASE HOMOLOG
9930	19401		2.38	2.0E-11	11417966	NT	Homo sapiens SEC14 (S. cerevisiae)-like 2 (SEC14L2), mRNA
650	10594	20412	1.24	1.0E-11	A 131016.1	NT	Homo sapiens SCL gene locus
1199	11109	20954	2.58	1.0E-11	AL 63279.2	NT	Homo sapiens chromosome 21 segment HS21C079
1483	11388		1.94	1.0E-11	AF 19914.1	NT	Homo sapiens PRO3078 mRNA, complete cds
2079	11989	21862	3.12	1.0E-11	AF000573.1	NT	Homo sapiens homogentisate 1,2-dioxygenase gene, complete cds
3454	13370	23174	1.32	1.0E-11	BE04315.1	EST HUMAN	CN0-BN0105-770300-292-d12 BN0105 Homo sapiens cDNA
5269	15191	24966	14.34	1.0E-11	AL 63247.2	NT	Homo sapiens chromosome 21 segment HS21C047
6773	16652	26840	3.25	1.0E-11	4885546	NT	Homo sapiens PHD finger protein 2 (PHF2) mRNA
6989	16896	27060	5.41	1.0E-11	R 13174.1	EST HUMAN	yJ73d08.s1 Seares infant brain 1NIH Homo sapiens cDNA clone IMAGE:28168 5'
7232	17109	27300	1.32	1.0E-11	BF 36519.1	EST HUMAN	QV4-NN1149-265090-423-a03 NN1149 Homo sapiens cDNA
7232	17109	27301	1.32	1.0E-11	BF 36519.1	EST HUMAN	QV4-NN1149-265090-423-a03 NN1149 Homo sapiens cDNA
8607	18474	28747	1.9	1.0E-11	BF 880078.1	EST HUMAN	602154807F1 NIH MGC_83 Homo sapiens cDNA clone IMAGE:4285977 5'
9702	19483		1.29	1.0E-11	ZZ 0377.1	EST HUMAN	HSAAACADH P_Human fetal Brain Whole tissue Homo sapiens cDNA
2922	12849	22649	0.82	9.0E-12	P 20742	SWISSPROT	PREGNANCY ZONE PROTEIN PRECURSOR (LSM34)
7641	17491	27712	1.22	9.0E-12	AL 63590.2	NT	Homo sapiens chromosome 21 segment HS21C100
7641	17491	27713	1.22	9.0E-12	AL 1163500.2	NT	Homo sapiens chromosome 21 segment HS21C100
9270	18974		3.57	8.0E-12	A 271736.1	NT	Homo sapiens Xq pseudoautosomal region; segment 2/2
4562	14454	24241	2.75	7.0E-12	Q 05904	SWISSPROT	34 KD SPICULIUM MATRIX PROTEIN PRECURSOR (LSM34)
8656	18545	28828	11.23	7.0E-12	AA 704735.1	EST HUMAN	zJ23d01.s1 Seares fetal liver_spleen_1NF1s_S1 Homo sapiens cDNA clone IMAGE:451152 3'
3500	13417		0.92	6.0E-12	AV 730554.1	EST HUMAN	AV730554 HTTF Homo sapiens cDNA clone HTTF/WFO6 5'
4252	14151	23925	7.85	6.0E-12	AA 732516.1	EST HUMAN	nz28f11.s1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1302573 3' similar to contains Alu repetitive element;
7201	17078	27263	1.19	6.0E-12	AF 003249.1	NT	Morone saxatilis myosin heavy chain FM3A (FM3A) mRNA, complete cds
7453	17262		1.86	6.0E-12	AA 847888.1	EST HUMAN	odf10g11.s1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1367588 similar to contains MER29.t2 MER29 repetitive element;
1027	10345	20790	2.88	5.0E-12	T 6573.1	EST HUMAN	EST04462_Fetal brain, Stratagene (cat#936206) Homo sapiens cDNA clone HFBDV33
3344	13264	23070	1.18	5.0E-12	BE 4779.1	EST HUMAN	Iz42b05.y1 NCI_CGAP_Bm62 Homo sapiens cDNA clone IMAGE:2291217 5'
3666	13580	23367	5.93	5.0E-12	A 271736.1	NT	Homo sapiens Xq pseudoautosomal region; segment 2/2
5171	15037		0.84	5.0E-12	AA 20661.1	EST HUMAN	nw24b11.s1 NCI_CGAP_GCB0 Homo sapiens cDNA clone IMAGE:1241373 3'
5667	15578	25676	4.75	5.0E-12	AL 63278.2	NT	Homo sapiens chromosome 21 segment HS21C078
5909	15815	25940	9.12	5.0E-12	AW 974760.1	EST HUMAN	EST386850 MAGE sequences, MAGN Homo sapiens cDNA
7268	17145	27359	2.15	5.0E-12	A 271735.1	NT	Homo sapiens Xq pseudoautosomal region; segment 1/2

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7895	17745			4.67	5.0E-12 AL163303.2	NT	Homo sapiens chromosome 21 segment HS2/C1'03
244	10211	20027		3.42	4.0E-12 AA700326.1	EST HUMAN	274g11.s1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:460676_3'
245	10211	20027		3.55	4.0E-12 AA700326.1	EST HUMAN	274g11.s1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:460676_3'
4620	14413	24198		0.85	4.0E-12 A1689984.1	EST HUMAN	b26105.x1 NCI CGAP_Lu24 Homo sapiens cDNA clone IMAGE:2270745_3 similar to TRQ13559 Q13559
6797	16676			2.89	4.0E-12 AF09907.1	NT	MARINER TRANSPOSEASE_;
8431	18305	28561		3.51	4.0E-12 A1229043.1	NT	Homo sapiens S164 gene, partial cds; PS1 and hypothetical protein genes, complete cds; and S171 gene, partial cds
9520	19131			1.9	4.0E-12 U78027.1	NT	Homo sapiens 959 kb contig between AML1 and CBR1 on chromosome 21q22, segment 3/3
600	10536	20345		3.81	3.0E-12 AW341683.1	EST HUMAN	Homo sapiens Bruton's tyrosine kinase (BTK), alpha-D-galactosidase A (GLA), L44-like ribosomal protein (L44L) and FTP3 (FTP3) genes, complete cds
600	10536	20346		3.81	3.0E-12 AW341683.1	EST HUMAN	hd13d01.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2909377_3' similar to TR-O14517 O14517 SMRP_;
8047	17938	28187		3.08	3.0E-12 U31672.1	NT	hd13d01.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2909377_3' similar to TR-O14517 O14517 SMRP_;
8047	17938	28188		3.08	3.0E-12 U31672.1	NT	Human prostate specific antigen gene, 5' flanking region
3421	13338	23143		1.03	2.0E-12 6754495	NT	Human prostate specific antigen gene, 5' flanking region
4025	13928	23704		1.04	2.0E-12 J01184.1	NT	Mus musculus keratin-associated protein 6.2 (Krtap6-2), mRNA
4025	13928	23705		1.04	2.0E-12 J01184.1	NT	Rat USA small nuclear RNA
4324	14221			1.8	2.0E-12 BE063509.1	EST HUMAN	Rat USA small nuclear RNA
4804	14688	24473		0.78	2.0E-12 O70306	SWISSPROT	CMD-BT0281-031199-087-a03 BT0281 Homo sapiens cDNA
4804	14688	24474		0.78	2.0E-12 O70306	SWISSPROT	TBX15 PROTEIN (T-BOX PROTEIN 15)
5893	15804			2.22	2.0E-12 AW971857.1	EST HUMAN	TBX15 PROTEIN (T-BOX PROTEIN 15)
6258	16124	26277		3.34	2.0E-12 T08169.1	EST HUMAN	EST38946 MAGE sequences, MAGL Homo sapiens cDNA
6516	16315	26552		2.07	2.0E-12 11422229	NT	EST38946 MAGE sequences, MAGL Homo sapiens cDNA
7367	17345			1.68	2.0E-12 AF1968664.1	NT	EST6060 Infant Brain, Bento Soares Homo sapiens cDNA clone HBBA13 5' end
7737	17587			8.13	2.0E-12 BE165990.1	EST HUMAN	Homo sapiens putative BPES syndrome breakpoint region protein gene, complete cds
9175	18916			1.71	2.0E-12 AL163283.2	NT	MR3-HT0487-150200-113-901 HT0487 Homo sapiens cDNA
9377	19041			1.52	2.0E-12 11418248	NT	Homo sapiens sulfotransferase-related protein (SULTX3), mRNA
117	10095	19914		1.82	1.0E-12 AW627674.1	EST HUMAN	hh90a09.x1 NCI CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2970040 3' similar to contains MER18.t1 MER18 repetitive element;
1944	11839			2.03	1.0E-12 A1871726.1	EST HUMAN	wm51077.x1 NCI CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2439493 3' similar to contains L1.b3 L1 repetitive element;
3032	12980	22752		1.16	1.0E-12 AF000991.1	NT	Homo sapiens testis-specific Testis Transcript Y 2 (TTY2) mRNA, partial cds

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
3032	12960	22753	1.16	1.0E-12 AF000891.1	NT	Homo sapiens testis-specific Testis Transcript Y2 (TTY2) mRNA, partial cds	
3798	13710	23496	27.7	1.0E-12 AU132248.1	EST_HUMAN	AU132248 NT2RP3 Homo sapiens cDNA clone NT2RP3004070 5'	
3798	13710	23497	27.7	1.0E-12 AU132248.1	EST_HUMAN	AU132248 NT2RP3 Homo sapiens cDNA clone NT2RP3004070 5'	
5630	15544		1.73	1.0E-12 U82828.1	NT	Homo sapiens ataxia telangiectasia (ATM) gene, complete cds	
5682	15591		1.82	1.0E-12 C9Y2G7	SWISSPROT	HYPOTHETICAL_ZINC FINGER PROTEIN_KIAA0861	
6224	16090	26240	1.72	1.0E-12 AF198864.1	NT	Homo sapiens putative BPES syndrome breakpoint region protein gene, complete cds	
6240	16106	26256	9.11	1.0E-12 A1248533.1	EST_HUMAN	qh66a04.x1 Soares_fetal_liver_spleen_1NF1S_S1 Homo sapiens cDNA clone IMAGE:11849614 3' similar to gb:M19503 LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (HUMAN); contains MER10 nt MER10 repetitive element;	
6240	16106	26257	9.11	1.0E-12 A1248533.1	EST_HUMAN	qh66a04.x1 Soares_fetal_liver_spleen_1NF1S_S1 Homo sapiens cDNA clone IMAGE:11849614 3' similar to gb:M19503 LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (HUMAN); contains MER10 nt MER10 repetitive element;	
7051	16928	27119	1.31	1.0E-12 AA782323.1	EST_HUMAN	ac26d05.s1 Stratagene ovary (#867217) Homo sapiens cDNA clone IMAGE:857577 3'	
9085	18880	29118	3.36	1.0E-12 AW962164.1	EST_HUMAN	EST374237 MAGE resequences, MAGG Homo sapiens cDNA	
9605	19728		2.2	1.0E-12 P44836	SWISSPROT	PROTABLE TONB-DEPENDENT RECEPTOR/H0712 PRECURSOR	
9963	19426		2.72	1.0E-12 X85949.1	NT	M.setulosus mitochondrial 12S rRNA gene	
3575	13489		1.13	9.0E-13 A1271735.1	NT	Homo sapiens Xq pseudoautosomal region; segment 1/2	
3865	13776	23570	1.16	9.0E-13 AE029890.1	NT	Homo sapiens CST gene for cerebroside sulfotransferase, exon 1, 2, 3, 4, 5	
7537	17288		2.37	9.0E-13 NE96653.1	EST_HUMAN	zg2606.s1 Soares_fetal_liver_spleen_1NF1S Homo sapiens cDNA clone IMAGE:293661 3'	
700	10633	20458	5.05	8.0E-13 U29185.1	NT	Homo sapiens prion protein (PrP) gene, complete cds	
700	10633	20459	5.05	8.0E-13 U29185.1	NT	Homo sapiens prion protein (PrP) gene, complete cds	
1796	11694	21570	1.36	8.0E-13 U80017.1	NT	Homo sapiens basic transcription factor 2 p44 (btf2p44) gene, partial cds, neuronal apoptosis inhibitory protein (nap1) and survival motor neuron protein (smn) genes, complete cds	
7838	17688		2.13	8.0E-13 U78027.1	NT	Homo sapiens Bruton's tyrosine kinase (BTK), alpha-D-galactosidase A (GLA), L44-like ribosomal protein (L44) and FTP3 (FTP3) genes, complete cds	
						Human germline T-cell receptor beta chain TCRBV13S1, TCRBV6S8A2T, TCRBV5SSA3N2T, TCRBV13SSA2T, TCRBV6SS9P, TCRBV5SS3A2T, TCRBV13S8P, TCRBV6S3A1N1T, TCRBV5S2, TCRBV6S6A2T, TCRBV5S7P, TCRBV13S4, TCRBV6S2A1N1T, TCRBV5S4A2T, TCRBV6S4A1,	
8984	18789	28079	2.49	8.0E-13 U66060.1	NT	TCRBV23S1A2T, TCRBV12-	
9550	19149		10.33	7.0E-13 BE778223.1	EST_HUMAN	601463285F1 NIH MGCG_67 Homo sapiens cDNA clone IMAGE:3866613 5'	
9768	19287		1.33	7.0E-13 Q10473	SWISSPROT	POLYPEPTIDE NACETYLGALACTOSAMINYLTRANSFERASE (PROTEIN-UDP-ACETYL GALACTOSAMINYL TRANSFERASE) (UDP-GALNAc:POLYPEPTIDE, N-ACETYL GALACTOSAMINYL TRANSFERASE) (GALNAc-T1)	
2054	11944	21840	18.51	6.0E-13 AL163207.2	NT	Homo sapiens chromosome 21 segment HS2/C007	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
3280	13201		1.05	5.0E-13	R78338.1	EST_HUMAN	yj82fc4.r1 Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE:145759 5'
3351	13271		1.45	5.0E-13	AA435773.1	EST_HUMAN	zt77a12.s1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:728350 3' similar to contains Alu repetitive element; contains element MER22 repetitive element;
8234	18115	28867	2.75	5.0E-13	P07213	SWISSPROT	MYO3IN LIGHT CHAIN KINASE, SKELETAL MUSCLE (MlCK)
1824	11721	-	2.96	4.0E-13	AW378614.1	EST_HUMAN	PM2+H70224-22_099-001-e11 HT0224 Homo sapiens cDNA
2411	12288		1.58	4.0E-13	AF003529.1	NT	Homo sapiens glyican 3 (GPC3) gene, partial cds and flanking repeat regions
5430	15350	25404	4.86	4.0E-13	BE169131.1	EST_HUMAN	PN3-H70520-23_020200-002-c08 HT0520 Homo sapiens cDNA
6278	16142	26298	1.92	4.0E-13	AB037750.1	NT	Homo sapiens mRNA for KIAA1329 protein, partial cds
6542	16400		1.52	4.0E-13	N44291.1	EST_HUMAN	yj63g05.r1 Soares melanocyte 2lbHM Homo sapiens cDNA clone IMAGE:273080 5' similar to PIR:AS2995 A32985 t complex sterility protein - mouse ;
7757	17607	27831	4.57	4.0E-13	AI288831.1	EST_HUMAN	qnr32d05.x1 NCI_CGAP_Kid5 Homo sapiens cDNA clone IMAGE:189894 5' similar to contains Alu repetitive element;
8507	18379	28645	1.83	4.0E-13	AA435819.1	EST_HUMAN	zt78g10.s1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:728514 3'
8507	18379	28646	1.83	4.0E-13	AA435819.1	EST_HUMAN	zt78g10.s1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:728514 3'
173	10144		3.8	3.0E-13	AF003528.1	NT	Homo sapiens X-linked arachnodactylic ectodermal dysplasia protein gene (EDA), exon 2 and flanking regions
847	10774		1.37	3.0E-13	AA430310.1	EST_HUMAN	zv68g08.r1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:781406 5'
2319	12200	22099	1.25	3.0E-13	AJ271736.1	NT	Homo sapiens X-linked pseudoautosomal region; segment 2/2
2429	12306		2.63	3.0E-13	AL163210.2	NT	Homo sapiens chromosome 21 segment HS21C010
2629	12497	22387	4.23	3.0E-13	BF372962.1	EST_HUMAN	CM3-F70100-14070-24270-08 F71010 Homo sapiens cDNA clone IMAGE:1324035 3'
3150	13075		2.86	3.0E-13	AA745844.1	EST_HUMAN	ob18dd2.s1 NCI_CGAP_Kid5 Homo sapiens cDNA clone IMAGE:1324035 3'
6601	16481	26669		3.0E-13	U52111.2	NT	Homo sapiens X28 region near ALD locus containing dual specificity phosphatase 9 (DUSP9), ribosomal protein L18a (RPL18a), Ca2+-Calmodulin-dependent protein kinase I (CAMK), creatine transporter (CRTTR), CDM protein (CDM), adrenoleukodystrophy protein >
8064	17955		4.03	3.0E-13	A1064768.1	EST_HUMAN	HA0536 Human fetal liver cDNA library Homo sapiens cDNA
8403	18279	28631	3.66	3.0E-13	BE063509.1	EST_HUMAN	CM0-B70281-031199-087-a03 BT0281 Homo sapiens cDNA
8876	18688	28980	2.6	3.0E-13	AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048
144	10118	19938	2.77	2.0E-13	U52111.2	NT	Homino sapiens X28 region near ALD locus containing dual specificity phosphatase 9 (DUSP9), ribosomal protein L18a (RPL18a), Ca2+/Calmodulin-dependent protein kinase I (CAMK), creatine transporter (CRTTR), CDM protein (CDM), adrenoleukodystrophy protein >
239	10207	20024	1.31	2.0E-13	U28839.1	NT	Danio rerio fibroblast growth factor receptor 4 mRNA, complete cds
1249	11156	21005	4.71	2.0E-13	AF239710.1	NT	Homo sapiens DNA polymerase delta small subunit (POLD2) gene, exons 1 through 11 and complete cds
3244	13167	22966	1.08	2.0E-13	BF431899.1	EST_HUMAN	hab7605.x1 Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE: 3'

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
4019	13923			1.72	2.0E-13 AL163278.2	NT	Homo sapiens chromosome 21 segment HS21C078
5722	15629	25732		3.87	2.0E-13 Q06852	SWISSPROT	CELL SURFACE GLYCOPROTEIN 1 PRECURSOR (OUTER LAYER PROTEIN B) (S-LAYER PROTEIN 1)
6078	16061	26209		6.32	2.0E-13 X16912.1	NT	Human PFKL gene for liver-type 6-phosphofructokinase (EC 2.7.1.11) exon 2
7976	17826	28067		3.97	2.0E-13 5031898	NT	Homo sapiens mab-21 (C. elegans)-like 1 (MAB21L1) mRNA
9251	18961			7.42	2.0E-13 AW892155.1	EST_HUMAN	CM0-NN0001-100300-274-e11 Nl00001 Homo sapiens cDNA
288	10252	20072		1.37	1.0E-13 S74129.1	NT	FGF-1-fibroblast growth factor 1 [human, kidney, Genomic, 342 nt, segment 2 of 2]
870	10736	20646		4.39	1.0E-13 AJ001973.1	NT	Homo sapiens LGMD2B gene
1314	11220	21077		1.27	1.0E-13 X87344.1	NT	H. sapiens DMA, DMB, HLA-Z1, IP2, LMP2, TAP1, LMP7, TAP2, DOB, DQB2 and RING8, 9, 13 and 14 genes
1976	11869	21761		2.16	1.0E-13 AA720574.1	EST_HUMAN	rw21g02.s1 NCI_CGAP_GCB0 Homo sapiens cDNA clone IMAGE:1241138 3' similar to contains THR.3
4488	14382	24169		1.48	1.0E-13 BF340987.1	EST_HUMAN	THR repetitive element; h0203809f1 NCI_CGAP_Bm64 Homo sapiens cDNA clone IMAGE:4185866 5'
8684	18872	28855		13.83	1.0E-13 BF108755.1	EST_HUMAN	7145s10.x1 Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE:3524443 3' similar to contains MER29,b2 MER29 repetitive element;
9076	18853			1.62	1.0E-13 AV715377.1	EST_HUMAN	AV715377 DCB Homo sapiens cDNA clone DCBAIE03 5'
9714	19255			1.6	1.0E-13 AJ271735.1	NT	Homo sapiens Xq pseudoautosomal region; segment 1/2
330	10289	20105		2.92	9.0E-14 AA781159.1	EST_HUMAN	aj24c01.s1 Soares_NHT Homo sapiens cDNA clone 1391232 3' similar to contains MER19;11 MER19 repetitive element;
331	10290	20106		2.85	9.0E-14 AA781159.1	EST_HUMAN	aj24c01.s1 Soares_NHT Homo sapiens cDNA clone 1391232 3' similar to contains MER19;11 MER19 repetitive element;
2451	12328			4.04	9.0E-14 AW861577.1	EST_HUMAN	RC4-CT0322-080100-013-d09 CT0322 Homo sapiens cDNA
2725	12587	22482		4.62	9.0E-14 AB038162.1	NT	Homo sapiens TFF gene cluster for trefoil factor, complete cds
3073	13000	22790		3.74	9.0E-14 AW513296.1	EST_HUMAN	xo54h05.x1 NCI_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2707833 3'
3200	10289	20105		0.98	9.0E-14 AA781159.1	EST_HUMAN	aj24c01.s1 Soares_NHT Homo sapiens cDNA clone 1391232 3' similar to contains MER19;11 MER19 repetitive element;
3728	13640	23426		5.22	9.0E-14 D14547.1	NT	Human DNA, SINE repetitive element
4660	14536	24325		1.93	9.0E-14 AJ002153.1	NT	Sequinu cedipus gene for seminal vesicle secreted protein semenogelin 1
3463	13369			1.57	8.0E-14 BE468263.1	EST_HUMAN	hz71cc9.x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3211324 3'
3872	13783			2.77	8.0E-14 RT6269.1	EST_HUMAN	yf172e03.11 Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE:144706 3'
7434	16447	26637		60.69	8.0E-14 X89211.1	NT	H. sapiens DNA for endogenous retroviral like element
7515	17303	27510		3.49	8.0E-14 AA219816.1	EST_HUMAN	zq17c0.s1 Strategene fetal retina 937202 Homo sapiens cDNA clone IMAGE:629970 3'
8732	18588			4.39	8.0E-14 BE062558.1	EST_HUMAN	QV2-BT0258-261098-014-a01 BT0258 Homo sapiens cDNA

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
1611	12699			3.07	7.0E-14 AW151673.1	EST_HUMAN	x6f1e10_x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2623146 3' similar to contains MER10_t2
363	10319	20140		10.2	6.0E-14 AF020503.1	NT	Homo sapiens FRA3B common fragile region, diadenosine triphosphate hydrolase (FHTT) gene, exon 5
5114	14982	24756		1.02	6.0E-14 8923548_NT	EST_HUMAN	Homo sapiens hypothetical protein FLJ20585 (FLJ20585), mRNA
5114	14982	24757		1.02	6.0E-14 8923548_NT	EST_HUMAN	Homo sapiens hypothetical protein FLJ20585 (FLJ20585), mRNA
7652	17502	27725		2.56	6.0E-14 AF020503.1	NT	Homo sapiens FRA3B common fragile region, diadenosine triphosphate hydrolase (FHTT) gene, exon 5
7652	17502	27726		2.56	6.0E-14 AF020503.1	NT	Homo sapiens FRA3B common fragile region, diadenosine triphosphate hydrolase (FHTT) gene, exon 5
602	10538	20348		3.92	5.0E-14 Q6S120	SWISSPROT	CANALICULAR MULTISPECIFIC ORGANIC ANION TRANSPORTER 1 (MULTIDRUG RESISTANCE ASSOCIATED PROTEIN 2) (CANALICULAR MULTIDRUG RESISTANCE PROTEIN)
4985	14860	24628		1.09	5.0E-14 AW073791.1	EST_HUMAN	x03605_x1 NCI_CGAP_GU1 Homo sapiens cDNA clone IMAGE:2575185 3' similar to contains L1_t2 L1
5397	15916	25393		5.12	5.0E-14 P08547	SWISSPROT	repetitive element;
1107	12685			1.77	4.0E-14 P04928	SWISSPROT	L1-REVERSE TRANSCRIPTASE HOMOLOG
1885	11732	21608		6.5	4.0E-14 AJ007973.1	EST_HUMAN	S-ANTIGEN PROTEIN PRECURSOR
3693	13607			0.94	4.0E-14 AA046502.1	EST_HUMAN	z67a06_r1 Scores_pregnant uterus_Nb6-HPU Homo sapiens cDNA clone IMAGE:487858 5'
4194	14094	23873		1	4.0E-14 N46328.1	EST_HUMAN	w73c12_s1 Scores_multiple_scenarios_2Nb6MSP Homo sapiens cDNA clone IMAGE:279190 3' similar to
9777	19760			2.31	4.0E-14 ABB86224.1	EST_HUMAN	contains L1_t3 L1 repetitive element;
934	10859	20705		2.13	3.0E-14 X95468.1	EST_HUMAN	wn08c03_x1 NCI_CGAP_Ut4 Homo sapiens cDNA clone IMAGE:2435532 3' similar to contains Alu
						NT	R.norvegicus mRNA for CPG2 protein
4841	14722	24505		0.82	3.0E-14 AW265354.1	EST_HUMAN	xp45f12_x1 NCI_CGAP_HN11 Homo sapiens cDNA clone IMAGE:2743343 3' similar to contains Alu
4844	14725	24507		1.1	3.0E-14 7656864_NT	EST_HUMAN	repetitive element; contains element MER9 repetitive element;
5104	14972			1.23	3.0E-14 BF466372.1	EST_HUMAN	Homo sapiens a disintegrin and metalloproteinase domain 29 (ADAM29), mRNA
5157	15024	24791		1.5	3.0E-14 P02894	SWISSPROT	hx94f11_x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:3195501 3' similar to contains MER4_b2
						EST_HUMAN	MER4 repetitive element;
						EST_HUMAN	CIRCUMSPOROZOITE PROTEIN PRECURSOR (CS)
8563	14722	24505		7.59	3.0E-14 AW265354.1	EST_HUMAN	xp45f12_x1 NCI_CGAP_HN11 Homo sapiens cDNA clone IMAGE:2743343 3' similar to contains Alu
384	10531	20154		3.98	2.0E-14 AJ274736.1	EST_HUMAN	repetitive element; contains element MER9 repetitive element;
384	10531	20155		3.98	2.0E-14 AJ274736.1	EST_HUMAN	Homo sapiens Xq pseudoautosomal region; segment 2/2
675	12673	20428		6.35	2.0E-14 AL165303.2	EST_HUMAN	Homo sapiens chromosome 21 segment HS21C103
						NT	Homo sapiens chromosomal region; segment 2/2

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2339	12219			1.36	2.0E-14 AW372888.1	EST_HUMAN	RC5-BT0377-091298-031-D12 BT0377 Homo sapiens cDNA
2416	12293			1.09	2.0E-14 7657539	NT	Homo sapiens rhabdoid tumor deletion region protein 1 (RTDR1), mRNA
2479	12355	22246		1.24	2.0E-14 AL163209.2	NT	Homo sapiens chromosome 21 segment HS21C009
2640	12507			0.95	2.0E-14 P08548	SWISSPROT	LIN-E-REVERSE TRANSCRIPTASE HOMOLOG
5505	15423	25485		2.96	2.0E-14 U01317.1	NT	Human beta globin region on chromosome 11
6114	16008			2.18	2.0E-14 BE0006550.1	EST_HUMAN	RC3-BN0072-24020-011-a06 BN0072 Homo sapiens cDNA
6432	16293	26454		19.91	2.0E-14 BE158761.1	EST_HUMAN	IL2-HT0397-071298-024-D04 HT0397 Homo sapiens cDNA
6432	16293	26455		19.91	2.0E-14 BE158761.1	EST_HUMAN	IL2-HT0397-071298-024-D04 HT0397 Homo sapiens cDNA
8160	18048	28300		4.76	2.0E-14 AW139800.1	EST_HUMAN	U1-H-B11-adw-a-10-0-J1_s1 NCL_CGAP_Sub3 Homo sapiens cDNA clone IMAGE:2718234 3'
8791	15423	25485		1.81	2.0E-14 U01317.1	NT	Human beta globin region on chromosome 11
1051	10968	20810		1.31	1.0E-14 AL163246.2	NT	Homo sapiens chromosome 21 segment HS21C046
1384	11289	21143		7.67	1.0E-14 AL163268.2	NT	Homo sapiens chromosome 21 segment HS21C068
1384	11289	21144		7.67	1.0E-14 AL163268.2	NT	Homo sapiens chromosome 21 segment HS21C068
1956	11851	21738		21.54	1.0E-14 L44140.1	NT	Homo sapiens chromosome X region from flamin (FLN) gene to glucose-6-phosphate dehydrogenase (G6PD) gene, complete cds's
2137	12025	21921		6.17	1.0E-14 AL163303.2	NT	Homo sapiens chromosome 21 segment HS21C103
2358	12238	22134		6.43	1.0E-14 AF001689.1	NT	Homo sapiens ribosomal protein L23A (RPL23A) gene, complete cds
2914	12841	22641		1.38	1.0E-14 P05227	SWISSPROT	HISTIDINE-RICH PROTEIN PRECURSOR (CLONE PFHRP-1)
3130	13055	22854		4.67	1.0E-14 BF335227.1	EST_HUMAN	RC2-CT0432-310700-013-a09_1 CT0432 Homo sapiens cDNA
3130	13055	22855		4.67	1.0E-14 BF335227.1	EST_HUMAN	RC2-CT0432-310700-013-a09_1 CT0432 Homo sapiens cDNA
3811	13723	23512		2	1.0E-14 AA682994.1	EST_HUMAN	aeg9c12.s1 Strategene schizo brain S11 Homo sapiens cDNA clone IMAGE:971350 3'
4374	14270	24051		1.74	1.0E-14 AW275852.1	EST_HUMAN	xk39n10.x1 NCL_CGAP_Lu28 Homo sapiens cDNA clone IMAGE:2753059 3'
5550	15466	25536		1.97	1.0E-14 AF126145.1	NT	Bos taurus xenobiotic/medium-chain fatty acid:CoA ligase form XL-III mRNA, nuclear mRNA encoding mitochondrial protein, complete cds
6012	19457	26045		10.41	1.0E-14 11487150	NT	Homo sapiens prionin (mouse)-like 1 (PROML1), mRNA
6012	19457	26046		10.41	1.0E-14 11487150	NT	Homo sapiens prionin (mouse)-like 1 (PROML1), mRNA
1558	11463	21320		2.85	9.0E-15 7427522	NT	Homo sapiens protein tyrosine phosphatase, receptor type, T (PTPRT), mRNA
2126	12014			1.64	9.0E-15 AF196779.1	NT	Homo sapiens transcription factor GHM enhancer 3, JM1 protein, JM5 protein, T54 protein, JM10 protein, A4 differentiation-dependent protein, triple LIM domain protein 6, and synaptophysin genes, complete cds; and L-type calcium channel $\alpha 2$
6427	16288	26449		4.28	9.0E-15 P21416	SWISSPROT	GAG POLYPROTEIN [CONTAINS: CORE PROTEINS P15, P12, P30, P10]
6673	16653	26748		1.53	9.0E-15 BE903595.1	EST_HUMAN	60177750F-NIH MGC_21 Homo sapiens cDNA clone IMAGE:3960156 5'
2780	10415			1	8.0E-15 BE261482.1	EST_HUMAN	601148632F-NIH MGC_19 Homo sapiens cDNA clone IMAGE:3164023 5'

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Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit No.	Top Hit Database Source	Top Hit Descriptor
7963	17813		2.83	7.0E-15 AW241958.1	EST_HUMAN	xn77d022x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone lMAGE:2700483 3' similar to contains THR12 THR repetitive element;	
978	10901	20748	6.12	6.0E-15 AJ271736.1	NT	Homo sapiens Xq pseudautosomal region; segment 2/2	
8622	19770		1.86	6.0E-15 AW836843.1	EST_HUMAN	QV1-L1T0036-150200-070-c10 L1T0036 Homo sapiens cDNA	
9948	19415		1.57	6.0E-15 BF432200.1	EST_HUMAN	hab81c12x1 Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone lMAGE:3' 3	
404	10350	20177	5.79	5.0E-15 AL163208.2	NT	Homo sapiens chromosome 21 segment HS21C008	
							Human hereditary haemochromatosis region, histone 2A-like protein gene, hereditary haemochromatosis (HLA-H) gene, RoRet gene, and sodium phosphate transporter (NPT3) gene, complete cds
2733	12595	22490	1.38	5.0E-15 Ug1328.1	NT	U1-H-BW0-alb-g-10-0-U1_NCL_CGAP_Sub6 Homo sapiens cDNA clone lMAGE:2731219 3'	
3423	13340		1	5.0E-15 AW296817.1	EST_HUMAN	AV730056 HTF Homo sapiens cDNA clone HTFAVE06 5'	
8063	17954		2.22	5.0E-15 AV730056.1	EST_HUMAN	Homo sapiens chromosome 21 segment HS21C103	
421	9988	19779	2.6	4.0E-15 AL163303.2	NT	DKFZp776I00810_R1_761 (synonym: ham2) Homo sapiens cDNA clone DKFZp761C0810 5'	
4039	13942	23720	0.78	4.0E-15 AL118596.1	EST_HUMAN	Homo sapiens mRNA for transcription factor	
8414	16438	26623	2.38	4.0E-15 AJ130894.1	NT	Homo sapiens mRNA for transcription factor	
8414	16438	26624	2.38	4.0E-15 AJ130894.1	NT	LY114ZF Human fetal heart, Lambda ZAP Express Homo sapiens cDNA clone LY114ZF 5' similar to ANFCARDIODILATIN)	
4123	14023		5.93	3.0E-15 N89452.1	EST_HUMAN	SWISSPROT NADH:UBIQUINONE OXIDOREDUCTASE CHAIN 5	
4843	14724		1.41	3.0E-15 P92485	EST_HUMAN	7P01Fc3 Chromosome 7 Placental cDNA Library Homo sapiens cDNA clone 7P01Fc3	
4955	14832	24589	0.88	3.0E-15 AA078097.1	EST_HUMAN	7P01Fc3 Chromosome 7 Placental cDNA Library Homo sapiens cDNA clone 7P01Fc3	
4955	14832	24600	0.88	3.0E-15 AA078097.1	EST_HUMAN	Mus musculus ultra high sulfur keratin gene, complete cds	
6314	16177	26535	2.86	3.0E-15 M27695.1	NT	Mus musculus ultra high sulfur keratin gene, complete cds	
6314	16177	26536	2.86	3.0E-15 M27695.1	NT	cc36a07_s1_NCL_CGAP_GCB1 Homo sapiens cDNA clone lMAGE:1351764 3' similar to contains MER19.11	
7709	17559		1.87	3.0E-15 AA807128.1	EST_HUMAN	MER19 repetitive element;	
8173	18061	28811	2.71	3.0E-15 AB026898.1	NT	Homo sapiens DNA, DLEC1 to ORC1L4 gene region, section 1/2 (DLEC1, ORC1L3, ORC1L4 genes, complete cds)	
250	10216	20033	3.29	2.0E-15 AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
364	10320	20141	3.23	2.0E-15 AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
364	10320	20142	3.23	2.0E-15 AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
1512	11417		1.14	2.0E-15 8923201	NT	Homo sapiens hypothetical protein FLJ20212 (FLJ20212), mRNA	

Table 4

Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
3465	13381	23186	1.04	2.0E-15 AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
3465	13381	23187	1.04	2.0E-15 AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
4522	14415		2.07	2.0E-15 AI806335.1	EST_HUMAN	wf0706.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2349923 3' similar to TR:Q61043 Q61043_NINEIN ;	
5097	14966	24741	1.33	2.0E-15 P13993	SWISSPROT	REPETITIVE PROLINE-RICH CELL WALL PROTEIN 2 PRECURSOR	
5097	14966	24742	1.33	2.0E-15 P13993	SWISSPROT	REPETITIVE PROLINE-RICH CELL WALL PROTEIN 2 PRECURSOR	
6223	16089		1.71	2.0E-15 AJ400877.1	NT	Homo sapiens ASCL3 gene, CEGP1 gene, C11orf14 gene, C11orf15 gene, C11orf16 gene and C11orf17 gene	
6310	16173	26631	2.2	2.0E-15 AA704195.1	EST_HUMAN	2J76d3.s1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:298675 5' similar to zt78d10.r1 Soares_fetal_lung_Ndrl-19W Homo sapiens cDNA clone IMAGE:460924 3'	
6375	16237	26397	5.13	2.0E-15 W505064.1	EST_HUMAN	WP:544f4.8 CE02227 TRANSPPOSEASE ;	
7163	17040	27232	2.72	2.0E-15 D14547.1	NT	Human DNA, SINE repetitive element	
7410	17277	27484	1.26	2.0E-15 AW379465.1	EST_HUMAN	CMD-HT0244-201099-078-a12 HT0244 Homo sapiens cDNA	
7410	17277	27485	1.26	2.0E-15 AW379465.1	EST_HUMAN	CMD-HT0244-201099-078-a12 HT0244 Homo sapiens cDNA	
8212	18096		3.01	2.0E-15 AJ271735.1	NT	Homo sapiens Xq pseudautosomal region; segment 1/2	
9799	13381	23186	2.22	2.0E-15 AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
9799	13381	23187	2.22	2.0E-15 AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
2747	12609		1.84	1.0E-15 AI89984.1	EST_HUMAN	fc26h05.x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:2270745 3' similar to TR:Q13539 Q13539 MARINER TRANSPOSEASE ;	
2979	12906	22706	0.8	1.0E-15 BE043584.1	EST_HUMAN	hk40502.y1 NCI CGAP_Ov34 Homo sapiens cDNA clone IMAGE:2999162 5'	
3103	13029	22825	0.99	1.0E-15 P08547	SWISSPROT	LINE-1 REVERSE TRANSCRIPTASE HOMOLOG	
4262	14161	23939	0.8	1.0E-15 BE182696.1	EST_HUMAN	RC3-HT0649-100500-022-h05 HT0649 Homo sapiens cDNA	
5086	14956	24731	1.15	1.0E-15 AI884928.1	EST_HUMAN	wr86g04.x1 NCI CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2494590 3'	
5844	15750	25864	1.83	1.0E-15 T95763.1	EST_HUMAN	ye40e10.s1 Soares_fetal_liver_spleen_1NFLS Homo sapiens cDNA clone IMAGE:120234 3' similar to contains MER6 repetitive element;	
6182	16068		1.96	1.0E-15 BE074217.1	EST_HUMAN	QV3-BT0569-270100-074-g05 BT0569 Homo sapiens cDNA	
6791	16670	26862	1.28	1.0E-15 AL163280.2	NT	Homo sapiens chromosome 21 segment HS21C080	
6905	16783	26976	4.57	1.0E-15 AI200976.1	EST_HUMAN	qf68h06.x1 Soares_nht Homo sapiens cDNA clone IMAGE:1755227 3'	
6905	16783	26977	4.57	1.0E-15 AI200976.1	EST_HUMAN	qf68h06.x1 Soares_nht Homo sapiens cDNA clone IMAGE:1755227 3'	
7227	17104	27293	1.44	1.0E-15	4507208 NT	Homo sapiens spermidine synthase (SRM) mRNA	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
8193	18079	28331	6.81	1.0E-15 AF044083.1	NT		Homo sapiens major histocompatibility locus class III region
9867	19492	25131	3.71	1.0E-15 AI783944.1	EST_HUMAN	tr31c05_x1_NCI_CGAP_Ov23 Homo sapiens cDNA clone IMAGE:2219912 3' similar to contains Alu repetitive element	
4404	14298	24082	1.03	9.0E-16	4503168 NT	Homo sapiens cut (Drosophila)-like 1 (CCAAT displacement protein) (CUTL1) mRNA	
8561	18238	28486	2.6	9.0E-16 F08688.1	EST_HUMAN	HSC22F05 normalized infant brain cDNA Homo sapiens cDNA clone c-2305	
6343	16206	26368	1.5	7.0E-16 Q88807	SWISSPROT	PROTEIN-ARGININE DEMINASE TYPE IV (PEPTIDYLARGININE DEMINASE IV) (PAD-R4)	
6343	16206	26369	1.5	7.0E-16 Q88807	SWISSPROT	PROTEIN-ARGININE DEMINASE TYPE IV (PEPTIDYLARGININE DEMINASE IV) (PAD-R4)	
9816	19585		6.8	7.0E-16 T94149.1	EST_HUMAN	ye28c12.1 Strategene lung (#937210) Homo sapiens cDNA clone IMAGE:119062 5'	
2094	11983		8.32	6.0E-16 AW972611.1	EST_HUMAN	EST384702 MAGE resequences, MAGL Homo sapiens cDNA	
1477	11382	21246	1.09	5.0E-16 AJ251154.1	NT	Mus musculus olfactory receptor cluster, OR37A, OR37B, OR37C, OR37E genes and OR37D pseudogene	
2047	12514	22404	1.79	5.0E-16 AA992176.1	EST_HUMAN	dt8004.51 Soares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:1623078 3' similar to	
7784	17634	27867	1.69	5.0E-16 AL163246.2	NT	contains element L1 repetitive element;	
8509	18623	28914	3.33	5.0E-16 BF217368.1	EST_HUMAN	601885734F1 NIH MGC_57 Homo sapiens cDNA clone IMAGE:4104129 5'	
9504	19381		8.34	5.0E-16	11418127 NT	Homo sapiens GTP binding protein 1 (GTPBP1) mRNA	
2192	12079		1.27	4.0E-16 AB001523.1	NT	Homo sapiens gene for TMEM1 and PWP2, complete and partial cds	
2828	12209	22107	1.32	4.0E-16 AW797168.1	EST_HUMAN	QV7-U0036-20030-115-q02 UN0036 Homo sapiens cDNA	
2828	12209	22108	1.32	4.0E-16 AW797168.1	EST_HUMAN	QV1-U0036-20030-115-q02 UN0036 Homo sapiens cDNA	
3411	13328	23129	3.85	4.0E-16 Q16653	SWISSPROT	MYELIN-OGLICODENDROCYTE GLYCOPROTEIN PRECURSOR	
4050	13952	23728	3.55	4.0E-16 BE083875.1	EST_HUMAN	PM4-BT0650-010400-002-009 BT0650 Homo sapiens cDNA	
4050	13952	23729	3.55	4.0E-16 BE083875.1	EST_HUMAN	PM4-BT0650-010400-002-009 BT0650 Homo sapiens cDNA	
5092	14962	24737	1	4.0E-16 P08548	SWISSPROT	LINE-1 REVERSE TRANSCRIPTASE HOMOLOG	
6538	16396	26375	33.8	4.0E-16 AL163284.2	NT	Homo sapiens chromosome 21 segment HS21C084	
7358	17226	27425	1.22	4.0E-16	11423191 NT	Homo sapiens hypothetical protein FLJ10024 (FLJ10024), mRNA	
8551	18421	28591	1.74	4.0E-16 AV730030.1	EST_HUMAN	AV730030 HTF Homo sapiens cDNA clone HTFWA03 5'	
9156	18906		1.64	4.0E-16 P08548	SWISSPROT	LINE-1 REVERSE TRANSCRIPTASE HOMOLOG	
9244	18957		5.94	4.0E-16 C05947.1	EST_HUMAN	C05947 Human pancreatic islet Homo sapiens cDNA clone hbc5355	
9255	18964	25519	2.04	4.0E-16	6912459 NT	Homo sapiens Grb2-associated binder 2 (KIAA0571), mRNA	
128	10102	19924	1.59	3.0E-16 AW022862.1	EST_HUMAN	df45c01.y1 Morton Fetal Cochlea Homo sapiens cDNA clone IMAGE:2486376 5'	
128	10102	19925	1.59	3.0E-16 AW022862.1	EST_HUMAN	df45c01.y1 Morton Fetal Cochlea Homo sapiens cDNA clone IMAGE:2486376 5'	
458	10402		1.47	3.0E-16 AL048445.1	EST_HUMAN	DKFZp43P037_11234 (synonym: hts3) Homo sapiens cDNA clone DKFZp43P037 5'	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
467	10410			1.6	3.0E-16 AF135446.1	NT	Homo sapiens TSX (TSX) pseudogene, exon 5
1436	11341	21207		2.01	3.0E-16 Q28983	SWISSPROT	ZONADIRESIN PRECURSOR
2046	12873	22670		4.05	3.0E-16 P03200	SWISSPROT	ENVELOPE GLYCOPROTEIN GP340 (MEMBRANE ANTIGEN) (MA) [CONTAINS: GLYCOPROTEIN GP220]
3874	13785			8.18	3.0E-16 AF020503.1	NT	Homo sapiens FRA3B common fragile region, diadenosine triphosphate hydrolase (FHTT) gene, exon 5
3875	13786			0.99	3.0E-16 U03887.1	NT	Human BXp20 gene
4861	14741	24521		1.01	3.0E-16 AV661393.1	EST_HUMAN	AV661393 GLC Homo sapiens cDNA clone GLCGSA01 3'
5452	15373	25431		1.41	3.0E-16 AF003529.1	NT	Homo sapiens glyican 3 (GPC3) gene, partial cds and flanking repeat regions
7031	16908	27098		4.72	3.0E-16 AI002886.1	EST_HUMAN	arm9h05.s1 Stratagene schizo brain S11 Homo sapiens cDNA clone IMAGE:1684185 3' similar to contains THR.b2 THR repetitive element;
7691	17541			1.27	3.0E-16 BF690817.1	EST_HUMAN	602246538F1 NIH MIGC_62 Homo sapiens cDNA clone IMAGE:43230322 5'
7828	17678	27922		3.08	3.0E-16 L78810.1	NT	Homo sapiens ADP/ATP carrier protein (ANT-2) gene, complete cds
956	10880			1.18	2.0E-16 AL163279.2	NT	Human chromosomes 21 segment HS21C079
2337	12217			0.96	2.0E-16 AA621761.1	EST_HUMAN	af0604.s1 Soares, testis_NHT Homo sapiens cDNA clone IMAGE:1030855 3'
2857	12524			1.71	2.0E-16 J03081.1	NT	Human SAAV-related endogenous retroviral LTR-like element
4087	13987	23764		1.33	2.0E-16 X89211.1	NT	H.sapiens DNA for endogenous retroviral like element
6646	16526	26720		1.63	2.0E-16 AF1732837.1	EST_HUMAN	n247f06.x5 NCI CGAP_Prt12 Homo sapiens cDNA clone IMAGE:1290947 similar to TR:054849 O54849 HYPOTHETICAL 42.9 KD PROTEIN. [2] TR:008905 ;contains MER7;[t] MER7 repetitive element;
178	10149	19963		2.55	1.0E-16 AF200719.1	NT	Homo sapiens pituitary tumor transforming gene protein (PTTG) gene, complete cds
377	10361			22.41	1.0E-16 AA628592.1	EST_HUMAN	af39g11.s1 Soares, total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:1034084 3' similar to contains OFR t2 OFR repetitive element;
1928	11823	21704		2.44	1.0E-16 BF327942.1	EST_HUMAN	QV0-BN0148-070700-295-a10 BN0148 Homo sapiens cDNA
5877	15783			23.72	1.0E-16 U45983.1	NT	Homo sapiens CCR8 chemokine receptor (CMKBR8) gene, complete cds
5960	15865	25987		2.81	1.0E-16 Q012779	SWISSPROT	MITOGEN-ACTIVATED PROTEIN KINASE KINASE 10 (MIXED LINEAGE KINASE 2) (PROTEIN KINASE MST)
6058	15783			6.59	1.0E-16 U45983.1	NT	Homo sapiens CCR8 chemokine receptor (CMKBR8) gene, complete cds
7349	17217	27416		1.31	1.0E-16 AW875651.1	EST_HUMAN	QV2-PT0012-040400-124-e05 PT0012 Homo sapiens cDNA
3679	13593	23379		2.54	9.0E-17 AW900048.1	EST_HUMAN	CM1-NN1003-200200-153-e01 NN1003 Homo sapiens cDNA
6035	15938			2.02	9.0E-17 A19392064.1	EST_HUMAN	Ig22c11-x1 NCI CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2109524 3' similar to contains MER28.t2 MER28 repetitive element;
6720	16600			4.87	9.0E-17 AW150257.1	EST_HUMAN	xg49j12.x1 NCI CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2630950 3' similar to contains OFR.t2 OFR repetitive element;

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
7870	117720		2.18	9.0E-17 AF200719.1	NT		Homo sapiens pituitary tumor transforming gene protein (PTTG) gene, complete cds
1001	10919		1.7	8.0E-17 AW380701.1	EST_HUMAN	QV0-OT0032-080300-155-d01 O7032 Homo sapiens cDNA	
3817	13729		0.87	8.0E-17 AL163280.2	NT	Homo sapiens chromosome 21 segment HS21C080	
5427	19444	25402	3.56	8.0E-17 BE172081.1	EST_HUMAN	MRO-HT0559-060300-003-e04 HT0559 Homo sapiens cDNA	
6311	16174		1.36	8.0E-17 AV730759.1	EST_HUMAN	AV730759 HTTF Homo sapiens cDNA clone HTFAQB07 5'	
1442	11347		3.18	7.0E-17 6753097	NT	Mus musculus apolipoprotein B editing complex 2 (Apobec2), mRNA	
5292	15184		3.05	7.0E-17 AF216650.1	NT	Homo sapiens putative MTAP (MTAP) mRNA, partial cds, alternatively spliced	
6017	15621	26052	6.83	7.0E-17 AF229843.1	NT		Mus musculus WNT-2 gene, partial cds; putative arylkinin-related protein and cystic fibrosis transmembrane conductance regulator (CFTR) genes, section 1 of 2 of the complete cds; and unknown gene
196	10168	19986	4.78	6.0E-17 AW9633880.1	EST_HUMAN	RC1-HN0003-220300-021-b04 HN0003 Homo sapiens cDNA hi81d04_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2978695 3' similar to contains h1.12	
5812	15717	25830	1.61	6.0E-17 AW662772.1	EST_HUMAN	L1 repetitive element :	
415	9882	19773	2.31	5.0E-17 T64110.1	EST_HUMAN	yc05m08_r1 Stratagene lung (##37210) Homo sapiens cDNA clone IMAGE:79839 5'	
6446	16335	26502	2.07	5.0E-17 T81048.1	EST_HUMAN	yd26b04_r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:109327 5'	
8783	18858	28887	2.12	4.0E-17 AL163247.2	NT	Homo sapiens chromosome 21 segment HS21C047	
9171	18914		1.98	4.0E-17 AI073546.1	EST_HUMAN	0145e04_x1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:1640286 3' similar to TR:Q16530	
2051	11941	21836	1.35	3.0E-17 AW119123.1	EST_HUMAN	Q16530_PMS3_MRNA;contains MER10.12 MER10 repetitive element ;	
3157	13082		1.31	3.0E-17 P38410	EST_HUMAN	xd89c09_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2604784 3'	
3590	13504	23293	1.14	3.0E-17 BE126522.1	EST_HUMAN	hw03b04_x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3181999 3'	
3590	13504	23294	1.14	3.0E-17 BE126522.1	EST_HUMAN	hw03b04_x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3181999 3'	
7594	17445	27660	4.72	3.0E-17 AB26888.1	NT	Homo sapiens DNA, DLEC1 to ORC1L4 gene region, section 1/2 (DLEC1, ORC1L3, ORC1L4 genes, complete cds)	
9134	18890		3.16	3.0E-17 11417966	NT	Homo sapiens SEC14 (S. cerevisiae)-like 2 (SEC14L2), mRNA	
9909	19386		15.82	3.0E-17 AV720204.1	EST_HUMAN	AV720204 GLC Homo sapiens cDNA clone GLCD1F08 5'	
350	10309	20127	2.81	2.0E-17 AI270080.1	EST_HUMAN	qf63ad6_x1 NCI_CGAP_Eso2 Homo sapiens cDNA clone IMAGE:1959922 3' similar to contains Alu repetitive element;	
351	10309	20127	2	2.0E-17 AI270080.1	EST_HUMAN	qf63ad6_x1 NCI_CGAP_Eso2 Homo sapiens cDNA clone IMAGE:1959922 3' similar to contains Alu repetitive element;	
972	10896		1.27	2.0E-17 AA722932.1	EST_HUMAN	zg81d04_s1 Soares fetal heart NbHHT9W Homo sapiens cDNA clone IMAGE:399751 3'	
2397	12276	22170	2.6	2.0E-17 Q23983	SWISSPROT	ZONADHESIN PRECURSOR	
2397	12275	22171	2.6	2.0E-17 Q23983	SWISSPROT	ZONADHESIN PRECURSOR	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
2899	12826	22621	5.62	2.0E-17 P12036	SWISSPROT	NEUROFILAMENT TRIPLETH PROTEIN (200 kDa NEUROFILAMENT PROTEIN) (NEUROFILAMENT HEAVY POLYPEPTIDE) (NF-H)	
5295	15216	25017	1.95	2.0E-17 M27685.1	NT	Mus musculus ultra high sulfur keratin gene, complete cds	
5295	15216	25018	1.95	2.0E-17 M27685.1	NT	Mus musculus ultra high sulfur keratin gene, complete cds	
5790	15636		2.07	2.0E-17 AF055066.1	NT	Homo sapiens MHC class 1 region	
6711	16591	26779	1.44	2.0E-17 Q95156	SWISSPROT	OLFACTOORY RECEPTOR-LIKE PROTEIN OLFR3	
6925	16803	26997	1.38	2.0E-17 AA300640.1	EST_HUMAN	EST13504 Testis tumor Homo sapiens cDNA 5' end similar to glycogenin	
7699	17549	27772	2.72	2.0E-17 AL163247.2	NT	Homo sapiens chromosome 21 segment HS21C047	
7699	17549	27773	2.72	2.0E-17 AL163247.2	NT	Homo sapiens chromosome 21 segment HS21C047	
7887	17737	27981	5.3	2.0E-17 D13391.1	NT	Human CYP19 genes for aromatase cytochrome P-450, promoter region (containing two cis-acting transcriptional regulatory elements)	
733	10695	20499	3.37	1.0E-17 P08183	SWISSPROT	MULTIDRUG RESISTANCE PROTEIN 1 (P-GLYCOPROTEIN 1)	
1729	11630	21499	2.45	1.0E-17 AL163207.2	NT	Homo sapiens chromosome 21 segment HS21C007	
2069	11959	21853	1.68	1.0E-17 P02461	SWISSPROT	COLLAGEN ALPHA ((III)) CHAIN PRECURSOR	
2287	12170	22057	1.46	1.0E-17 U79410.1	NT	Homo sapiens thrombospondin 2 (THBS2) gene, promoter region and exons 1A and 1B	
3519	13435		1.01	1.0E-17 AF224669.1	NT	Homo sapiens mannosidase, beta A, lysosomal (MANBA) gene, and ubiquitin-conjugating enzyme E2D 3 (UBE2D3) genes, complete cds	
4045	13947		7.17	1.0E-17 R05942.1	EST_HUMAN	y30ed7_r1 Soares fetal liver spleen 1NF1S Homo sapiens cDNA clone IMAGE:1263888 5'	
5885	15792		4.7	1.0E-17 AW468468.1	EST_HUMAN	hex8ed5_x1 NCI CGAP_CML1 Homo sapiens cDNA clone IMAGE:2921312 3' similar to contains Alu repetitive element; contains LTR8.1L LTR8 repetitive element;	
6006	15911	26037	1.44	1.0E-17 A1185642.1	EST_HUMAN	qee5b65_x1 Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone IMAGE:1743825 3'	
6006	15911	26038	1.44	1.0E-17 A1185642.1	EST_HUMAN	qee5b65_x1 Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone IMAGE:1743825 3'	
6216	16082	26232	1.32	1.0E-17 Q16831	SWISSPROT	URIDINE PHOSPHORYLASE (UDRPASE)	
8719	18536	28821	2.01	1.0E-17 Q28824	SWISSPROT	MYOSIN LIGHT CHAIN KINASE, SMOOTH MUSCLE (MLCK) [CONTAINS: TELOKIN]	
2422	12289	22196	0.96	9.0E-18 AA174078.1	EST_HUMAN	zpf18g12_s1 Strategene fetal retina 937202 Homo sapiens cDNA clone IMAGE:609862 3'	
7468	17328		3.26	9.0E-18 AI472167.1	EST_HUMAN	jj8ed03_x1 Soares_NSFF8_9W_OT_PA_P_SuH Homo sapiens cDNA clone IMAGE:2143889 3'	
3718	13650	23415	1.75	8.0E-18	4758977 NT	Homo sapiens protein tyrosine phosphatase, non-receptor type substrate 1 (PTPN1) mRNA	
346	10305	20121	8.39	7.0E-18 AW316976.1	EST_HUMAN	xx10b04_x1 NCI CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2837071 3' similar to gbl20868 60S RIBOSOMAL PROTEIN L4 (HUMAN);	
346	10305	20122	8.39	7.0E-18 AW316976.1	EST_HUMAN	xx10b04_x1 NCI CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2837071 3' similar to gbl20868 60S RIBOSOMAL PROTEIN L4 (HUMAN);	
5136	15003	24774	0.85	7.0E-18 R16220.1	EST_HUMAN	ya9ed07_s1 Soares infant brain 1NIB Homo sapiens cDNA clone IMAGE:653285 3' similar to contains L1 repetitive element;	

Table 4
Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
9638	10305	20121	5.73	7.0E-18 AW316976.1	EST_HUMAN	xx10b04_x1_NCL_CGAP_Pari Homo sapiens cDNA clone IMAGE:2837071 3' similar to gb:L208668 60S RIBOSOMAL PROTEIN L4 (HUMAN);	
9638	10305	20122	5.73	7.0E-18 AW316976.1	EST_HUMAN	xx10b04_x1_NCL_CGAP_Pari Homo sapiens cDNA clone IMAGE:2837071 3' similar to gb:L208668 60S RIBOSOMAL PROTEIN L4 (HUMAN);	
3255	13178	22976	1.16	6.0E-18 X71791.2	NT	Rattus norvegicus partial Gdnf/Pn-1 gene for glia-derived neitin/protease nexin 1, enhancer region PROTEIN-GLUTAMINE GAMMA-GLUTAMYL TRANSFERASE (TISSUE TRANSGLUTAMINASE)	
4641	14529		3.37	6.0E-18 P52181	SWISSPROT	(TGASE C) (TGC)	
6799	16678		2.69	6.0E-18	11428155	NT	Homo sapiens similar to high-mobility group (nonhistone chromosomal) protein 4 (H. sapiens) (LOC63446), mRNA
8476	18349	28614	1.75	6.0E-18 AL163246.2	NT	Homo sapiens chromosome 21 segment HS21C046	
8642	18506	28784	1.78	6.0E-18 X87344.1	NT	Homo sapiens D1MA, D1MB, HLA-Z1, IPP2, LMP2, TAP1, LMP7, TAP2, DOB, DQB2 and RING8, 9, 13 and 14 genes	
9394	19052	25308	3.66	6.0E-18 U87929.1	NT	Human acconitate hydratase (ACO2) gene, exon 4 qm65g11_x1 Scares_placenta_8tc9weeks_2NbHPsite&W Homo sapiens cDNA clone IMAGE:1893668 3'	
1130	11044	20886	14.74	5.0E-18 A1280214.1	EST_HUMAN	similar to contains Alu repetitive element;	
4217	14115	23893	0.92	5.0E-18	10946665	NT	Mus musculus galectinin (Gsdm), mRNA
5058	14928	24700	1.76	5.0E-18 D61517.1	EST_HUMAN	HUM411F05B Clontech human fetal brain polyA+ mRNA (#6335) Homo sapiens cDNA clone GEN-411F05	
5224	15147	24914	1.38	5.0E-18 AF087913.1	NT	5'	
7061	16938	27128	4.26	5.0E-18 BE143312.1	EST_HUMAN	Human endogenous retrovirus HERV-P-T47D MR0-HT0161-221098-002-c06 HT0161 Homo sapiens cDNA	
8346	18223	28474	4.33	5.0E-18	10242378	NT	Homo sapiens lymphocyte activation-associated protein (LOC51088), mRNA
8346	18223	28475	4.33	5.0E-18	10242378	NT	Homo sapiens lymphocyte activation-associated protein (LOC51088), mRNA
9512	19125		5.3	5.0E-18 AW867182.1	EST_HUMAN	MR1-SNC035-050400-001-g11 SN0035 Homo sapiens cDNA	
9837	19336		13	5.0E-18 AV860547.1	EST_HUMAN	AV650547 GLC Homo sapiens cDNA clone GI_CCGA023	
119	10096	19915	1.36	4.0E-18 BE044076.1	EST_HUMAN	ho36104_x1_NCL_CGAP_Urt Homo sapiens cDNA clone IMAGE:3039511 3' similar to contains MER29,b3 MER29 repetitive element;	
119	10096	19916	1.36	4.0E-18 BE044076.1	EST_HUMAN	ho36104_x1_NCL_CGAP_Urt Homo sapiens cDNA clone IMAGE:3039511 3' similar to contains MER29,b3 MER29 repetitive element;	
1846	11742		1.03	4.0E-18 A1738392.1	EST_HUMAN	wi33h08_x1_NCL_CGAP_Cor16 Homo sapiens cDNA clone IMAGE:2392095 3'	
2154	12042	21940	0.98	4.0E-18 Q06430	SWISSPROT	N-ACETYLGLUCOSAMINYLTRANSFERASE (N-BRANCHING ENZYME) (IGNT)	
2154	12042	21941	0.98	4.0E-18 Q06430	SWISSPROT	N-ACETYLGLUCOSAMINYLTRANSFERASE (N-BRANCHING ENZYME) (IGNT)	
5293	15214	25014	2.55	4.0E-18 A1017565.1	EST_HUMAN	ou23e06_x1 Scares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:1627138 3'	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Acession No.	Top Hit Database Source	Top Hit Descriptor
5263 15214	25015	2.55	4.0E-18	AI0177565.1	EST_HUMAN	ot23e06.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:1627138 3'	
8370 18247	28499	7.12	4.0E-18	AA371807.1	EST_HUMAN	EST83633 Pituitary gland, subtracted (prolactin/growth hormone) II Homo sapiens cDNA 5' end similar to EST containing O family repeat	
831 10758	20609	2.38	3.0E-18	AA814196.1	EST_HUMAN	ch23h11.s1 NCI_CGAP_Kid5 Homo sapiens cDNA clone IMAGE:1324581 3' similar to SW:RS5_HUMAN P46782_40S RIBOSOMAL PROTEIN S5.;	
914 10338	20686	2.41	3.0E-18	BE088634.1	EST_HUMAN	CM0-BT0690-210300-298-g07 BT0690 Homo sapiens cDNA	
3867 13778	23571	1.19	3.0E-18	AL163247.2	NT	Homo sapiens chromosome 21 segment HS21C047	
6084 16029	26169	5.2	3.0E-18	BE01671.1	EST_HUMAN	PM0-BN0081-100300-001-b08 BN0081 Homo sapiens cDNA	
9642 19210		4.92	3.0E-18	AW022015.1	EST_HUMAN	df31h12.y1 Morton Fetal Cochlea Homo sapiens cDNA clone IMAGE:2485126 5'	
251 10217	20034	2.83	2.0E-18	AW836820.1	EST_HUMAN	QV1-L-T0036-150200-070-e07 LT0036 Homo sapiens cDNA	
1135 11049		47.22	2.0E-18	BE25097.1	EST_HUMAN	6011142552F1 NIH_MGC_16 Homo sapiens cDNA clone IMAGE:3355044 5'	
5322 15242		3.2	2.0E-18	AA488610.1	EST_HUMAN	ak63sa07.x1 Soares_festis_NIH Homo sapiens cDNA clone IMAGE:1409652 3' similar to TR:O14577	
5380 15298	25149	3.04	2.0E-18	D14547.1	NT	O14577 BAC CLONE RG114A06 FROM 7231, COMPLETE SEQUENCE. ;	
5380 15298	25150	3.04	2.0E-18	D14547.1	NT	Human DNA, SINE repetitive element	
5585 15500		1.67	2.0E-18	BF347229.1	EST_HUMAN	6020211f4F1 NCI_CGAP_Bm67 Homo sapiens cDNA clone IMAGE:4156670 5'	
5814 15720	25834	3.53	2.0E-18	AW665853.1	EST_HUMAN	h194g01.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2979984 3' similar to contains MER19.12 MER19 repetitive element;	
7779 17629	27861	1.53	2.0E-18	AW151673.1	EST_HUMAN	xf67e10x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2623146 3' similar to contains MER10.12 MER10 repetitive element;	
7779 17629	27862	1.53	2.0E-18	AW151673.1	EST_HUMAN	xf67e10x1 NCI_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2623146 3' similar to contains MER10.12 MER10 repetitive element;	
8340 18217	28469	5.32	2.0E-18	AW470791.1	EST_HUMAN	ha33d06.x1 NCI_CGAP_Kid12 Homo sapiens cDNA clone IMAGE:2623146 3' similar to contains THR.b3	
8968 18774	29065	4.44	2.0E-18	AW151289.1	EST_HUMAN	THR repetitive element; xg47eq9.x1 NCI_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2630728 3' similar to contains MER8.b2	
9325 11049		3.15	2.0E-18	BE256097.1	EST_HUMAN	MER8 repetitive element; ha33d06.x1 NCI_CGAP_Kid12 Homo sapiens cDNA clone IMAGE:2623146 3' similar to contains THR.b3	
4318 14215		1.02	1.0E-18	T95406.1	EST_HUMAN	ye43g05.r1 Soares fetal liver spleen INFLS Homo sapiens cDNA clone IMAGE:120536 5' similar to contains L1 repetitive element;	
5286 15208	24985	2.38	1.0E-18	AV653405.1	EST_HUMAN	AV653405 GLC Homo sapiens cDNA clone GLODKE11 3'	
5419 16340	25594	1.97	1.0E-18	D0099.1	NT	Homo sapiens mRNA for Na,K-ATPase alpha-subunit, complete cds	
5419 16340	25595	1.97	1.0E-18	D0099.1	NT	Homo sapiens mRNA for Na,K-ATPase alpha-subunit, complete cds	
5883 15790	25912	1.32	1.0E-18	AL163280.2	NT	Homo sapiens chromosome 21 segment HS21C030	

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Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor	
7696	17546	27769	4.4	1.0E-18	U91328.1	NT	Human hereditary haemochromatosis region, histone 2A-like protein gene, hereditary haemochromatosis (HLA-H) gene, RoRet gene, and sodium phosphate transporter (NP73) gene, complete cds	
9277	18980	28324	2.53	1.0E-18	AF003529.1	NT	Homo sapiens glyican 3 (GPC3) gene, partial cds and flanking repeat regions	
533	10475	20289	4.28	9.0E-19	AA231961.1	EST_HUMAN	ztf1d06.r1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:712811 5' similar to contains MER19.12 MER19 repetitive element;	
534	10475	20289	3.19	9.0E-19	AA231961.1	EST_HUMAN	ztf1d06.r1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:712811 5' similar to contains MER19.12 MER19 repetitive element;	
6584	16464	16464	6.21	9.0E-19	F05688.1	EST_HUMAN	HSC23F051 normalized infant brain cDNA Homo sapiens cDNA clone c-23f05	
7042	16919	27110	2.46	9.0E-19	AL163203.2	NT	Homo sapiens chromosome 21 segment HS21C003	
7042	16919	27111	2.46	9.0E-19	AL163203.2	NT	Homo sapiens chromosome 21 segment HS21C003	
8470	18343	28608	3.83	9.0E-19	AB032969.1	NT	Homo sapiens mRNA for KIAA1143 protein, partial cds	
9042	10475	20289	8.69	9.0E-19	AA231961.1	EST_HUMAN	ztf1d06.r1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:712811 5' similar to contains MER19.12 MER19 repetitive element;	
1032	10950		1.54	8.0E-19	AW974902.1	EST_HUMAN	EST387007 MAGE sequences, MAGN Homo sapiens cDNA	
2198	12085	21987	1.58	7.0E-19	4758139	NT	Homo sapiens DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 6 (RNA helicase, 54kD) (DDX6) mRNA	
5884	15791	28913	2.15	7.0E-19	AF002090.1	NT	Rattus norvegicus cpt151 mRNA, partial cds	
9179	19742		2.95	7.0E-19	AA705684.1	EST_HUMAN	ZB60b01.s1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:435145 3'	
3713	13626		1.02	6.0E-19	AW852860.1	EST_HUMAN	ZB60b01.s1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:435145 3' PM0-CT0248-(31098-001-901 CT0248 Homo sapiens cDNA	
4360	14256	24041	1.36	6.0E-19	P34986	SWISSPROT	OLFACTOORY RECEPTOR 6 (M50)	
4360	14256	24042	1.36	6.0E-19	P34986	SWISSPROT	OLFACTOORY RECEPTOR 6 (M50)	
4694	14580		1.16	6.0E-19	AJ211795.1	NT	Homo sapiens Xq_pseudoautosomal region; segment 1/2	
4952	14829	24595	1.09	6.0E-19	AL120817.1	EST_HUMAN	DKEZp762F192_r1 762 (synonym: hme12) Homo sapiens cDNA clone DKEZp762F192_5'	
5571	15486	25562	5.24	5.0E-19	Q00193	SWISSPROT	ZONA PELLUCIDA SFERM-BINDING PROTEIN B PRECURSOR (ZONA PELLUCIDA GLYCOPROTEIN ZP-X) (RC55)	
8824	18637	28921	7.19	5.0E-19	AW183725.1	EST_HUMAN	x87602.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2664171 3' similar to contains element M5R1 repetitive element;	
							Human germline T-cell receptor beta chain TCRBV13S1, TCRBV6S8A2T, TCRBV5S6A3N2T, TCRBV13S6A2T, TCRBV6S9P, TCRBVSS3A2T, TCRBV13S8P, TCRBV6S3A1N1T, TCRBV5S2, TCRBV6S6A2T, TCRBV5S7P, TCRBV13S4, TCRBV6S2A1N1T, TCRBV5S4A2T, TCRBV6S4A1, TCRBV23S1A2T, TCRBV13S1A2T, TCRBV13S1V12>	
9849	19495			1.52	5.0E-19	U666060.1	NT	Homo sapiens mRNA, chromosome 1 specific transcript KIAA0501
542	10483	20293	1.45	4.0E-19	AB007970.1	NT	602130910F1 NIH_MGCG_56 Homo sapiens cDNA clone IMAGE:287674 5'	
2649	12516	22406	1.39	4.0E-19	BF697362.1	EST_HUMAN		

Table 4

Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
3778	13690	23475		3.0E-19 Q28997	SWISSPROT	BETA-2 ADRENERGIC RECEPTOR	
3778	13690	23476		3.0E-19 Q28997	SWISSPROT	BETA-2 ADRENERGIC RECEPTOR	
4345	14242	24024		3.0E-19 AV708136.1	EST_HUMAN	AV708136 ADC Homo sapiens cDNA clone ADCAmA11 5'	
6370	16232			2.47	3.0E-19 11432214	NT	Homo sapiens similar to aldo-keto reductase family 1, member B11 (aldo-keto reductase-like) (H. sapiens) (LOC63222), mRNA
7443	16456	26645		1.23	3.0E-19 X89685.1	NT	M_musculus mRNA for TPCR33 protein
9416	19068			7.38	3.0E-19 AF165520.1	NT	Homo sapiens phorbolin I protein (PBI) mRNA, complete cds
2513	12387	22279		17.57	2.0E-19 AL163201.2	NT	Homo sapiens chromosome 21 segment HS21C001
6849	16728	26923		8.96	2.0E-19 AA012854.1	EST_HUMAN	ze24cd8_r1 Soares retina N2b4H1R Homo sapiens cDNA clone IMAGE:3606880 5'
473	10417			1.6	1.0E-19 BE408611.1	EST_HUMAN	601304/125F1NIH_MGC_21 Homo sapiens cDNA clone IMAGE:36368310 5'
2118	12007	21906		1.4	1.0E-19 H30795.1	EST_HUMAN	yo79g07_r1 Soares adult brain N2b4H56Y Homo sapiens cDNA clone IMAGE:184188 5' similar to contains MER10 repetitive element;
2685	12550			2.88	1.0E-19 D38044.1	NT	Human gene for Ah-receptor, exon 7-9
2817	12746			5.03	1.0E-19 4758977	NT	Homo sapiens protein tyrosine phosphatase, non-receptor type substrate 1 (PTPN51) mRNA
3355	13274	23075		1.27	1.0E-19 AA834967.1	EST_HUMAN	aj49b12_s1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:1393651 3' similar to contains MER27_t2
5701	15609	25711		2.37	1.0E-19 U1286.1	NT	Oryctocephalus clinicus sodium/dicarboxylate cotransporter mRNA, partial cds
6919	16797	26990		1.79	1.0E-19 M64657.1	NT	Rabbit phosphorylase kinase beta subunit mRNA, complete cds
7076	16953			2.83	1.0E-19 T99620.1	EST_HUMAN	ye72b02_r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:123243 5' similar to contains MER37 repetitive element;
7853	17703	27948		25.21	1.0E-19 AW812259.1	EST_HUMAN	RC0-ST0174-181098-031-b05 ST0174 Homo sapiens cDNA
7857	17707	27953		1.79	1.0E-19 N44631.1	EST_HUMAN	y31e09_r1 Soares melanocyte 2NbHM Homo sapiens cDNA clone IMAGE:272872 5'
6005	15910	26035		2.22	8.0E-20 7657286	NT	Mus musculus keratin-associated protein 9-1 (Krtap9-1), mRNA
6005	15910	26036		2.22	8.0E-20 7657286	NT	Mus musculus keratin-associated protein 9-1 (Krtap9-1), mRNA
6439	16300	26462		1.31	8.0E-20 A122137.1	EST_HUMAN	q986f109_r1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:1842089 3'
6439	16300	26463		1.31	8.0E-20 A122137.1	EST_HUMAN	q986f109_r1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:1842089 3'
3238	13161	22261		0.88	7.0E-20 BF326455.1	EST_HUMAN	PM4-AN0096-050900-003-e04 AN0096 Homo sapiens cDNA
6176	15133	24852		5.53	7.0E-20 AL138120.1	EST_HUMAN	DKFZp547D092_r1 547 (synonym: hfr1) Homo sapiens cDNA clone DKFZp547D092 5'
6947	16825	27018		10.99	7.0E-20 AA557657.1	EST_HUMAN	MER29 repetitive element;
6947	16825	27019		10.99	7.0E-20 AA557657.1	EST_HUMAN	nl46c04_s1 NCL_CGAP_Pt4 Homo sapiens cDNA clone IMAGE:1043718 similar to contains MER29_b2
8952	18759			10.45	7.0E-20 6912633	NT	Homo sapiens ribosomal protein L13a (RPL13A), mRNA
3508	13424	23227		3.85	6.0E-20 P39188	SWISSPROT	ALU SUBFAMILY J SEQUENCE CONTAMINATION WARNING ENTRY

Table 4

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
4175	14075	23850	2.55	6.0E-20	BE22434.1	EST_HUMAN	601441231f1 NIH_MGC_72 Homo sapiens cDNA clone IMAGE:3916231 5'
4491	14385		1.17	5.0E-20	AV725123	EST_HUMAN	AV725123 HTC Homo sapiens cDNA clone HTCBTA01 5'
6634	16514	26704	4.66	5.0E-20	W90525.1	EST_HUMAN	z178d08.s1 Soares_fetal_liver_spleen_1NF1S_S1 Homo sapiens cDNA clone IMAGE:418191 3' similar to contains MER30.1t MER30 repetitive element;
6634	16514	26705	4.66	5.0E-20	W90525.1	EST_HUMAN	z178d08.s1 Soares_fetal_liver_spleen_1NF1S_S1 Homo sapiens cDNA clone IMAGE:418191 3' similar to contains MER30.1t MER30 repetitive element;
7128	17005	27197	1.44	5.0E-20	AB028174.1	NT	Mus musculus MMANG mRNA, complete cds
7128	17005	27198	1.44	5.0E-20	AB028174.1	NT	Mus musculus MMANG mRNA, complete cds
6624	16504		6.03	4.0E-20	Al874352.1	EST_HUMAN	z16af03.x1 NCI_CGAP_Ot35 Homo sapiens cDNA clone IMAGE:2293396 3'
7985	17845	28087	1.17	4.0E-20	AW987469.1	EST_HUMAN	QV3-DT0043-0902020-080-004 DT0043 Homo sapiens cDNA
2092	11981	21876	1.04	3.0E-20	U03888.1	NT	Human BXp21 gene
4115	14015	23795	1.69	3.0E-20	P23273	SWISSPROT	OLFACTORY RECEPTOR-LIKE PROTEIN 14
4524	14417	24201	0.86	3.0E-20	AA037616.1	EST_HUMAN	z136b12.s1 Soares_pregnant uterus_NbHPU Homo sapiens cDNA clone IMAGE:484895 3' similar to
7172	17049		3.26	3.0E-20	D14547.1	NT	contains L1.3 L1 repetitive element;
						Human DNA, SINE repetitive element	
8055	17946		2.08	3.0E-20	P11369	SWISSPROT	RETROVIRUS-RELATED POLYPOLYPROTEIN [CONTAINS: REVERSE TRANSCRIPTASE ; ENDONUCLEASE]
9195	18929	25552	5.37	3.0E-20	BE883422.1	EST_HUMAN	601514180f1 NIH_MGC_71 Homo sapiens cDNA clone IMAGE:3915522 5'
813	10741		4.52	2.0E-20	AW303868.1	EST_HUMAN	x24e10.x1 NCI_CGAP_U4 Homo sapiens cDNA clone IMAGE:2761098 3' similar to SW.RS5_MOUSE
1095	11011	20852	2.03	2.0E-20	AA516335.1	EST_HUMAN	ng69n109.s1 NCI_CGAP_Lip2 Homo sapiens cDNA clone IMAGE:9400097 similar to TR:G1224066
1095	11011	20853	2.03	2.0E-20	AA516335.1	EST_HUMAN	ng69n109.s1 NCI_CGAP_Lip2 Homo sapiens cDNA clone IMAGE:9400097 similar to TR:G1224066
2786	10741		2.72	2.0E-20	AW303868.1	EST_HUMAN	x24e10.x1 NCI_CGAP_U4 Homo sapiens cDNA clone IMAGE:2761098 3' similar to SW.RS5_MOUSE
4866	14746	24525	4.32	2.0E-20	Q28983	SWISSPROT	P97461_40S RIBOSOMAL PROTEIN S5. ;
4866	14746	24526	4.32	2.0E-20	Q28983	SWISSPROT	ZONADHESIN PRECURSOR
5090	14960		11.35	2.0E-20	5174538	NT	Homo sapiens malate dehydrogenase 1, NAD (soluble) (MDH1) mRNA
7301	17177	27378	2.95	2.0E-20	D10083.1	NT	Homo sapiens RGH1 gene, retrovirus-like element
7301	17177	27379	2.95	2.0E-20	D10083.1	NT	Homo sapiens RGH1 gene, retrovirus-like element
8993	18797	29089	1.95	2.0E-20	AA766755.1	EST_HUMAN	o835b08.s1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1306935 3' similar to contains MER4.b2 MER4 repetitive element;

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Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
8993	18797	29090		1.95	2.0E-20 AA769755.1	EST_HUMAN	oa35b08_s1 NCL_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1306935 3' similar to contains MER4_b2 MER4 repetitive element;
9574	19482	25129		1.5	2.0E-20 H55371.1	EST_HUMAN	CHR220310 Chromosome 22 exon Homo sapiens cDNA clone C22_391 5' zt1d06_s1 NCL_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:712811 5' similar to contains MER19_t2 MER19 repetitive element;
1967	12648	21751		5.25	1.0E-20 AA281961.1	EST_HUMAN	hr84b06_x1 NCL_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:3135155 3' similar to contains L1_t2_L1 repetitive element;
4938	14235	24018		1	1.0E-20 BF115158.1	EST_HUMAN	Homo sapiens Autosomal Highly Conserved Protein (AHCP), mRNA
7285	17161	27559		2.42	1.0E-20 11418491_NT		Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced
8839	18652	28940		2.81	1.0E-20 AF223391.1	NT	nc80g08_s1 NCL_CGAP_Pri1 Homo sapiens cDNA clone IMAGE:745694 similar to contains L1_t3_L1 repetitive element;
9321	19011			1.84	1.0E-20 AA420453.1	EST_HUMAN	AJ00354 Selected chromosome 21 cDNA library Homo sapiens cDNA clone MP1p12-8-21
2883	12810			1.08	9.0E-21 A1003514.1	EST_HUMAN	RC3-NN0068-09050-021-603 NN0068 Homo sapiens cDNA
9045	18830			2.35	9.0E-21 AW898189.1	EST_HUMAN	bb60a02_s1 NH ₂ -MGC_10 Homo sapiens cDNA clone IMAGE:2984714 5' similar to SW:NIAM_HUMAN
7115	16992			1.74	8.0E-21 AW674891.1	EST_HUMAN	O95169_NADH-UBIQUINONE OXIDOREDUCTASE ASH1 SUBUNIT PRECURSOR;
8825	18658	28922		3.42	8.0E-21 AA4809411.1	EST_HUMAN	ob71f06_s1 NCL_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1336835 3'
9207	18938			4.02	8.0E-21 O21330	SWISSPROT	ATP SYNTHASE A CHAIN (PROTEIN 6)
2023	11914	21803		2.07	7.0E-21 P15800	SWISSPROT	LAMININ BETA-2 CHAIN PRECURSOR (SLAMMIN) (LAMININ CHAIN B3)
2023	11914	21804		2.07	7.0E-21 P15800	SWISSPROT	LAMININ BETA-2 CHAIN PRECURSOR (SLAMMIN) (LAMININ CHAIN B3)
4161	14061			4.96	7.0E-21 AA046502.1	EST_HUMAN	z467ad6_s1 Soares_pregnant uterus NbHPU Homo sapiens cDNA clone IMAGE:4878568 5'
6887	16766	26962		1.43	7.0E-21 AJ277557.1	NT	Homo sapiens dNT-2 gene for mitochondrial 5'(3')-deoxyribonucleotidase (dNT-2 gene), exons 1-5
7036	16913	27102		6.84	7.0E-21 D14718.1	NT	Human chromosomal protein HMGI related gene
							zg75d03_s1 Soares_fetal_heart_NbHH19W Homo sapiens cDNA clone IMAGE:398981 3' similar to qb.M14338 VITAMIN K-DEPENDENT PROTEIN S PRECURSOR (HUMAN);contains THR,t3 OFR repetitive element;
8080	17971	28220		2.94	7.0E-21 AA723404.1	EST_HUMAN	Hom sapiens PTD013 protein (PTD013), mRNA
8590	18458	28727		2.37	7.0E-21 770668_NT		601304125F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:3638310 5'
4014	13920	23696		0.94	6.0E-21 BE408611.1	EST_HUMAN	Hom sapiens protein tyrosine phosphatase, non-receptor type 21 (PTPN21), mRNA
907	10831	20678		1.6	5.0E-21 5902031_NT		om23g03_s1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:1541908 3' similar to TR:O02711
2234	12119	22021		1.08	5.0E-21 AA928194.1	EST_HUMAN	002711 PRO-POL-DUTPASE POLYPROTEIN;
4266	14165	23942		2.65	5.0E-21 BE9658839.1	EST_HUMAN	601619871F1 NIH_MGC_74 Homo sapiens cDNA clone IMAGE:3933880 5'
4696	14582	24375		5.98	5.0E-21 4885474_NT		Homo sapiens melanoma antigen, family C, 1 (MAGEC1), mRNA

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
5109	149777			0.95	5.0E-21 D87675.1	NT	Homo sapiens DNA for amyloid precursor protein, complete cds
1701	11602	21473		1.24	4.0E-21 AA970713.1	EST_HUMAN	co886e08_s1 NCI_CGAP_Kid5 Homo sapiens cDNA clone IMAGE:1573094 3' similar to TR:Q16530 Q16530 PM53 mRNA; contains OFFR_tf OR repetitive element;
6106	16000	26138		3.06	4.0E-21 AB019576.1	NT	Rattus norvegicus mRNA for TIM, complete cds
2228	12113	22015		1.06	3.0E-21 AL163201.2	NT	Homo sapiens chromosome 21 segment HS21C001
3041	12968	22762		4.04	3.0E-21 AJ007973.1	NT	Homo sapiens LGMD2B gene
5749	15657			1.72	3.0E-21 BF184739.1	EST_HUMAN	601844465F1 NIH_MGC_54 Homo sapiens cDNA clone IMAGE:4064945 5'
6206	15966	28100		4.79	3.0E-21 BF361053.1	EST_HUMAN	RC1-O70083-100800-019-q08 OT0083 Homo sapiens cDNA
7589	17440	27656		1.55	3.0E-21 AW897760.1	EST_HUMAN	OMI-NIN063-280400-203-h08 NN0063 Homo sapiens cDNA
9684	19672	24992		1.32	3.0E-21 AL163213.2	NT	Homo sapiens chromosome 21 segment HS21C013
140	10114			14.75	2.0E-21 BE163247.1	EST_HUMAN	QY3-HT0458-170200-090-912 HT0458 Homo sapiens cDNA
919	10843	20688		0.85	2.0E-21 AB007857.2	NT	Homo sapiens mRNA for KIAA0397 protein, partial cds
919	10843	20689		0.85	2.0E-21 AB007857.2	NT	Homo sapiens mRNA for KIAA0397 protein, partial cds
1196	11106			2.09	2.0E-21 BE064410.1	EST_HUMAN	RC4-BT0311-141199-011-h06 BT0311 Homo sapiens cDNA
2599	12468	22361		2.89	2.0E-21 Q28983	SWISSPROT	ZONADHESIN PRECURSOR
2599	12468	22362		2.89	2.0E-21 Q28983	SWISSPROT	ZONADHESIN PRECURSOR
5370	15290	25126		1.77	2.0E-21 A1624582.1	EST_HUMAN	ts30f03_x1 NCI_CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2230109 3' similar to TR:Q99854 Q99854
6870	16749	26944		4.66	2.0E-21 BE141795.1	EST_HUMAN	HYPOTHETICAL_51_1_KD PROTEIN;
7122	16898	27190		3.43	2.0E-21 AU136779.1	EST_HUMAN	QV0-HT0103-091199-050-911 HT0103 Homo sapiens cDNA
8412	18287			1.98	2.0E-21 BE250127.1	EST_HUMAN	AU136779 PLACE1 Homo sapiens cDNA clone PLACE1005052 5'
8636	18501	28776		1.92	2.0E-21 BE973829.1	EST_HUMAN	ht09g01_x1 NCI_CGAP_Kid13 Homo sapiens cDNA clone IMAGE:3146256 3' similar to contains MER29b3
8636	18501	28777		1.92	2.0E-21 BE973829.1	EST_HUMAN	MER29 repetitive element;
9425	19072			5.13	2.0E-21 AF176815.1	NT	601680636F1 NIH_MGC_83 Homo sapiens cDNA clone IMAGE:3951008 5'
1235	11142	20994		1.68	1.0E-21 AA557657.1	EST_HUMAN	Hom sapiens putative 8-hydroxyguanine DNA glycosylase gene, complete cds
1381	11286			2.46	1.0E-21 AI601264.1	EST_HUMAN	ht4604_s1 NCI_CGAP_P74 Homo sapiens cDNA clone IMAGE:1043718 similar to contains MER29b2
5907	15813			2.43	1.0E-21 AL079752.1	EST_HUMAN	MER29 repetitive element;
6269	16134	26289		4.43	1.0E-21 AI223104.1	EST_HUMAN	ar88d12_x1 Barsleed colon HPLRB1 Homo sapiens cDNA clone IMAGE:2152343 3'
8021	17871			1.46	1.0E-21 57300383	NT	DKFZp734/058_J1_434 (synonym: hess3) Homo sapiens cDNA clone DKFZp734/058 5'
4312	14209	23993		2.55	9.0E-22 AI702438.1	EST_HUMAN	qq47e05_x1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:1638338 3' similar to gb:M64241 QM
							PROTEIN (HUMAN);
							Homo sapiens SET domain and maniner transposase fusion gene (SETMAR) mRNA
							I294403_x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2296204 3' similar to TR:Q15408 Q15408 NEUTRAL PROTEASE LARGE SUBUNIT;

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit No.	Top Hit Database Source	Top Hit Descriptor
7002	16879	27070		1.26	9.0E-22	AL163201.2	NT
7002	16879	27071		1.26	9.0E-22	AL163201.2	NT
8171	18059	28309		3.65	9.0E-22	AV761874.1	EST_HUMAN
8945	18753	29048		2.92	9.0E-22	AU140358.1	EST_HUMAN
8997	18800			2.39	9.0E-22	9790236	NT
933	10858			4.55	8.0E-22	BE144748.1	EST_HUMAN
6609	16489			3.8	8.0E-22	KA046502.1	EST_HUMAN
648	10585	20401		4.23	7.0E-22	AL163246.2	NT
4186	14086	23861		2.32	7.0E-22	Q611838	SWISSPROT
4969	14844	24613		1.18	7.0E-22	AE008881.1	NT
7044	16921			1.34	7.0E-22	AF151054.1	NT
7127	17004	27196		2.63	7.0E-22	M78590.1	EST_HUMAN
7538	17389	27599		1.86	7.0E-22	AF009860.1	NT
6796	16675			1.82	6.0E-22	AV029123.1	EST_HUMAN
5927	15632	25955		2.53	5.0E-22	AL163303.2	NT
7912	17762	28001		3.97	5.0E-22	U60822.1	NT
9645	19213			2.03	5.0E-22	BF476511.1	EST_HUMAN
3584	13498			0.85	4.0E-22	AL1271735.1	NT
6302	19767			2.7	4.0E-22	AL163202.2	NT
8105	17995	28244		2.97	4.0E-22	BF218030.1	EST_HUMAN
9803	19215			1.51	4.0E-22	AL163209.2	NT
943	10868			1.2	3.0E-22	AI469679.1	EST_HUMAN
2523	12297	22288		1.86	3.0E-22	AI859038.1	EST_HUMAN
3620	13534			1.49	3.0E-22	D14718.1	NT
4685	14581	24374		2.72	3.0E-22	AI090125.1	EST_HUMAN
6793	16672	26894		2.75	3.0E-22	BE089841.1	EST_HUMAN
1910	11805			2.86	2.0E-22	N24942.1	EST_HUMAN
2476	12352	22244		1.33	2.0E-22	P24916	SWISSPROT
3373	13292	23091		4.06	2.0E-22	8394043	NT
4133	14033	23808		1.26	2.0E-22	AV817794.1	EST_HUMAN
						PM1-ST0262-261199-001-d12-S1262	Homo sapiens cDNA

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5570	19447	25561		1.44	2.0E-22 W39456.1	EST_HUMAN	zc20f01.r1 Soares_senescent_fibroblasts_NbHSF Homo sapiens cDNA clone IMAGE:3222873 5' similar to gb:X72208 MONOCYTE CHEMOTACTIC PROTEIN 3 PRECURSOR (HUMAN);
5747	15655	25763		3.39	2.0E-22 BF092116.1	EST_HUMAN	RC0-TN0079-150900-025-h12 TN0079 Homo sapiens cDNA qJ76n06.x1 Soares_NhhMPu_S1 Homo sapiens cDNA clone IMAGE:1878299 3' similar to contains MER29 i3 MER29 repetitive element;
7595	17446	27661		1.49	2.0E-22 A1276522.1	EST_HUMAN	nt04h11.51 NCI_CGAP_Pt22 Homo sapiens cDNA clone IMAGE:1219268 3'
7640	17490	27710		7.07	2.0E-22 AA715315.1	EST_HUMAN	nt04h11.51 NCI_CGAP_Pt22 Homo sapiens cDNA clone IMAGE:1219269 3'
7640	17490	27711		7.07	2.0E-22 AA715315.1	EST_HUMAN	nt04h11.51 NCI_CGAP_Pt22 Homo sapiens cDNA clone IMAGE:1219269 3'
8974	18779	29071		2.04	2.0E-22 AW418960.1	EST_HUMAN	ha24f04.x1 NCI_CGAP_Kid12 Homo sapiens cDNA clone IMAGE:2874655 3'
9009	19304	25200		1.85	2.0E-22 AL163280.2	NT	Homo sapiens chromosome 21 segment HS21C080
1836	11733	21609		1.78	1.0E-22 AW865517.1	EST_HUMAN	PM4-SN0020-010400-009-h02 SN0020 Homo sapiens cDNA
2539	12413	22303		2.65	1.0E-22 U50871.1	NT	Human familial Alzheimer's disease (STM2) gene, complete cds
3365	13284	23084		1.74	1.0E-22 D14547.1	NT	Human DNA, SINE repetitive element
6553	16411	26589		1.49	1.0E-22 BE084667.1	EST_HUMAN	M0-B10639-22020-002-h07 B10659 Homo sapiens cDNA
9844	19342			4.71	9.0E-23 AW802801.1	EST_HUMAN	IL2-UM0076-070400-061-F11 UM0076 Homo sapiens cDNA
3523	13459	23236		0.84	8.0E-23 AF1983479.1	NT	Gallus gallus Dach2 protein (Dach2) mRNA, complete cds
5241	15165	24935		1.48	8.0E-23 AI133716.1	EST_HUMAN	HA2340 Human fetal liver cDNA library Homo sapiens cDNA
5241	15165	24936		1.48	8.0E-23 AI133716.1	EST_HUMAN	HA2340 Human fetal liver cDNA library Homo sapiens cDNA
3271	13192			1.45	7.0E-23 AV647246.1	EST_HUMAN	AV647246 GLC Homo sapiens cDNA clone GLCAW C07 3'
4764	14649	24438		1.04	7.0E-23 10092626	NT	Homo sapiens DKFZP756400463 protein (DKFZP756400463) mRNA
8398	18274	28526		3.51	7.0E-23 5031952	NT	Homo sapiens Not56 (D. melanogaster)-like protein (NOT56) mRNA
3387	13305			1.62	6.0E-23 AF199333.1	NT	Rattus norvegicus RIMB (Rim1B) mRNA, complete cds
4171	14071	23846		3.12	6.0E-23 AL163249.2	NT	Homo sapiens chromosome 21 segment HS21C049
9146	18898	25338		1.99	6.0E-23 AF224669.1	NT	Homo sapiens mannosidase, beta A, lysosomal (MANBA) gene, and ubiquitin-conjugating enzyme E2D 3 (UBE2D3) genes, complete cds
9146	18899	25339		1.99	6.0E-23 AF224669.1	NT	Homo sapiens mannosidase, beta A, lysosomal (MANBA) gene, and ubiquitin-conjugating enzyme E2D 3 (UBE2D3) genes, complete cds
9341	19021	25297		2.18	6.0E-23 AI209130.1	EST_HUMAN	qg56c03.x1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:1889460 3' similar to SW-MV10_MOUSE F23249 PROTEIN MOV-10 ;
5341	15262	25088		3.51	5.0E-23 U82671.2	NT	Homo sapiens chromosome Xq28 melanoma antigen family A2a (MAGEA2A), melanoma antigen family A2b (MAGEA2B), melanoma antigen family A3 (MAGEA3), cathepsin (CALT), NAD(P)H dehydrogenase-like protein (NSDH1), and L1 >
5781	19454	25797		3.51	5.0E-23 AF179818.1	NT	Pongo pygmaeus olfactory receptor (FPY116) gene, partial cds
6397	19454	25797		3.04	5.0E-23 AF179818.1	NT	Pongo pygmaeus olfactory receptor (FPY116) gene, partial cds

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
65580	16460	26653	3.56	3.0E-23 AA130165.1	EST_HUMAN	Z35g09..1 Soares_pregnant_uterus_NbHPU Homo sapiens cDNA clone IMAGE:503968 5' similar to	
7329	17233	27435	3.61	3.0E-23 Z70664.1	NT	contains MER29.i2 MER29 repetitive element;	
7329	17233	27436	3.61	3.0E-23 Z70664.1	NT	Human endogenous retroviral element HC2	
650	10586	20402	4.36	2.0E-23 AJ288880.1	NT	Human endogenous retroviral element HC2	
1126	12644		2.77	2.0E-23 M55270.1	NT	Homo sapiens KIAA0851 gene (partial), X73 gene and Z7FL1 gene	
2765	12627	22519	1.06	2.0E-23 P22105	SWISSPROT	Human matrix Gla protein (MGP) gene, complete cds	
2765	12627	22620	1.06	2.0E-23 P22105	SWISSPROT	TENASCIN-X PRECURSOR (TNX) (HEXBRACTION-LIKE)	
3325	13245		1.46	2.0E-23 AJ21458.1	EST_HUMAN	TENASCIN-X PRECURSOR (TNX) (HEXBRACTION-LIKE)	
3855	13569		3.03	2.0E-23 BE165980.1	EST_HUMAN	qS7311..X1 NCL CGAP_Prc28 Homo sapiens cDNA clone IMAGE:1943757 3' similar to TR:Q13537 Q13537	
3894	13804	23589	2.98	2.0E-23 H59931.1	EST_HUMAN	MER37 TRANSPOSABLE ELEMENT, COMPLETE CONSENSUS SEQUENCE. ;	
3894	13804	23590	2.98	2.0E-23 H59931.1	EST_HUMAN	MR3-HT0487-150200-113-g01 HT0487 Homo sapiens cDNA	
						Y716a02..1 Soares fetal liver spleen INFSL Homo sapiens cDNA clone IMAGE:205418 5'	
6595	16475		5.62	2.0E-23 AF280107.1	NT	Y716a02..1 Soares fetal liver spleen INFSL Homo sapiens cDNA clone IMAGE:205418 5'	
9131	18888		2.69	2.0E-23 M32658.1	NT	Homo sapiens cytochrome P450 polypeptide 43 (CYP3A43) gene, partial cds; cytochrome P450 polypeptide 4 (CYP3A4) and cytochrome P450 polypeptide 7 (CYP3A7) genes, complete cds; and cytochrome P450 polypeptide 5 (CYP3A5) gene, partial cds	
9656	19218		2.47	2.0E-23 AF009660.1	NT	Human alcohol dehydrogenase gamma subunit (ADH3) gene, exon 1	
9774	19676		2.02	2.0E-23 AU133831.1	EST_HUMAN	Home sapiens T cell receptor beta locus, TCRBV7/3A2 to TCRBV12S2 region	
4428	14323	24110	1.6	1.0E-23 AL165252.2	NT	AUJ33931 OVARC1 Homo sapiens cDNA clone OVARC1000946 5'	
4657	14543		4.49	1.0E-23 AL165210.2	NT	Homologous chromosome 21 segment HS21C052	
6034	15937		2.91	1.0E-23 BE373471.1	EST_HUMAN	Homologous chromosome 21 segment HS21C010	
						6012364..55F1 NIH_MGC_44 Homo sapiens cDNA clone IMAGE:36086653 5'	
6865	16744	26937	4.54	1.0E-23 AA448097.1	EST_HUMAN	ZW82C06..r1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:782698 5' similar to contains PTR5..2	
540	10481		1.88	9.0E-24 AA653213.1	EST_HUMAN	PTR5 repetitive element;	
4549	14442	24225	1.08	8.0E-24 P23269	SWISSPROT	ab75a08..s1 Strategene fetal retina S37202 Homo sapiens cDNA clone IMAGE:852758 3' similar to	
4549	14442	24226	1.08	8.0E-24 P23269	SWISSPROT	TR: E19822 E19822 CA PROTEIN ;	
3796	13708		1.31	7.0E-24 AW937954.1	EST_HUMAN	OLFACTOORY RECEPTOR-LIKE PROTEIN 13	
690	10623		2.4	6.0E-24 AB001421.1	NT	QV0-DT0047-170200-122-016 D10047 Homo sapiens cDNA	
820	10748	20595	10.14	6.0E-24 AL163249.2	NT	Macaca fuscata mRNA for Testis-Specific Protein Y (TSPY), complete cds	
3889	13800	23985	7.18	5.0E-24 AJ229043.1	NT	Home sapiens chromosome 21 segment HS21C049	
5611	15526		3.06	4.0E-24 AA594178.1	EST_HUMAN	Homologous 959 kb contig between AMI.1 and CBR1 on chromosome 21(q22, segment 3/3	
						mn311h05..s1 NCL CGAP_Gas1 Homo sapiens cDNA clone IMAGE:1085529 3' similar to SW:POL_MLVRK	
						F31795 POL POLYPROTEIN;	

Table 4
Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
9506	19121	25293		2.12	4.0E-24 AB029016.1	NT	Homo sapiens mRNA for KIAA1093 protein, partial cds
9748	19309	25203		1.53	4.0E-24	11418318 NT	Homo sapiens G-2 and S-phase expressed 1 (GTS1), mRNA
							hh68c08.s1 NCI CGAP GU1 Homo sapiens cDNA clone IMAGE:2967950 3' similar to contains MER29.b2
6906	16784			3.02	3.0E-24 AW614871.1	EST HUMAN	MER29 repetitive element;
7449	17258	27463		4.12	3.0E-24 AL163252.2	NT	Homo sapiens chromosome 21 segment HS21C052
9587	19171	25273		5.14	3.0E-24 BF127762.1	EST HUMAN	601810449F1 NIH MGIC_46 Homo sapiens cDNA clone IMAGE:4053396 5'
2298	12180	22078		2.33	2.0E-24 AA167539.1	EST HUMAN	zp11f09.r1 Stratagene fetal retina 937202 Homo sapiens cDNA clone IMAGE:6009161 5'
3729	13641			1.01	2.0E-24 AW898189.1	EST HUMAN	RC3-NN0068-980500-021-h03 NN0068 Homo sapiens cDNA
7075	16952	27146		3.14	2.0E-24 AL119158.1	EST HUMAN	DKFZp761L1712_r1 761 (synonym: hamy2) Homo sapiens cDNA clone DKFZp761L1712 5'
9433	19717			6.55	2.0E-24 M28877.1	NT	Human O family dispersed repeat element
1670	11572	21438		2.18	1.0E-24	77063340 NT	Homo sapiens CG-127 protein (LOC5i646), mRNA
2638	12505			1.87	1.0E-24 AW820194.1	EST HUMAN	QNU-ST0294-10400-185-c10 ST0294 Homo sapiens cDNA
2987	12915	22710		0.91	1.0E-24 D86423.1	NT	Mus musculus mRNA for HG11 keratin, partial cds
4173	14073			1.63	1.0E-24 AF433313.1	NT	Homo sapiens PTEN (PTEN) gene, exon 2
6456	16317	26484		4.07	1.0E-24 AL163303.2	NT	Homo sapiens chromosome 21 segment HS21C103
6633	16513	26703		1.98	1.0E-24 AW901164.1	EST HUMAN	CM0-NN010-130300-281-d07 NN1010 Homo sapiens cDNA
4926	14805	24575		2.32	7.0E-25 AA483944.1	EST HUMAN	ne92e10.s1 NCI CGAP Kid1 Homo sapiens cDNA clone IMAGE:911754 similar to contains MER1.b2
6785	16664	26855		3.75	7.0E-25 AA468646.1	EST HUMAN	MER1 repetitive element;
8942	18750	29045		7.46	7.0E-25 AA583540.1	EST HUMAN	nf25h06.s1 NCI CGAP_Prl Homo sapiens cDNA clone IMAGE:914843 similar to SW:R14A_YEAST
6174	15131			4.32	6.0E-25 W87623.1	EST HUMAN	P36105 PROBABLE 60S RIBOSOMAL PROTEIN L14A.;
6543	16401	26580		11.44	6.0E-25	7305360 NT	EST_HUMAN
5166	15032	24799		0.95	5.0E-25	AW838171.1	EST_HUMAN
8634	18499	28774		3.61	5.0E-25	AW979107.1	EST_HUMAN
1430	11335	21201		2.08	4.0E-25	T98107.1	EST_HUMAN
3356	13275			2.78	4.0E-25	AW887671.1	EST_HUMAN
4221	14119			3.02	4.0E-25	BE170957.1	EST_HUMAN
3218	13199	22989		2.66	3.0E-25	8923321 NT	Hom sapiens hypothetical protein FLJ20344 (FLJ20344), mRNA
3218	13199	23000		2.66	3.0E-25	8923321 NT	Hom sapiens hypothetical protein FLJ20344 (FLJ20344), mRNA
4798	14683	24470		0.85	3.0E-25	P29622 SWISSPROT	KALLISTATIN PRECURSOR (KALLIKREIN INHIBITOR) (PROTEASE INHIBITOR 4)
6854	16733	26926		2.47	3.0E-25	AL1163210.2	NT
							Homo sapiens chromosome 21 segment HS21C010

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
8394	18270	28522	2.03	3.0E-25 AA579013.1	EST_HUMAN	nt30h10.s1 NCL_CGAP_P11 Homo sapiens cDNA clone MAGE:915331 similar to contains L1.f1 L1	
1325	11232	21088	3.37	2.0E-25	5032158 NT	Homo sapiens transducin (beta)-like 1 (TBL1) mRNA	
2280	12144	22043	7.11	2.0E-25 BE888016.1	EST_HUMAN	601511530f1 NIH MGC_71 Homo sapiens cDNA clone MAGE:3913087 5'	
2801	12370	22293	4.32	2.0E-25 P17008	SWISSPROT	40S RIBOSOMAL PROTEIN S16	
4096	13996	23773	1.91	2.0E-25 P17008	SWISSPROT	40S RIBOSOMAL PROTEIN S16	
4096	13996	23774	1.91	2.0E-25 P17008	SWISSPROT	40S RIBOSOMAL PROTEIN S16	
7629	17480	27700	2.25	2.0E-25 AL449573.1	EST_HUMAN	AL449573 Homo sapiens Testis (Stavrides GS) Homo sapiens cDNA	
361	10317	20138	1.61	1.0E-25 AL040229.1	EST_HUMAN	DKFZp434H0373_r1 434 (synonym: htes3) Homo sapiens cDNA clone DKFZp434H0373 5'	
1228	11136		1.21	1.0E-25	9655487 NT	Human endogenous retrovirus, complete genome	
2384	12264	22156	2.79	1.0E-25 Q06055	SWISSPROT	ATP SYNTHASE LIPID-BINDING PROTEIN P2 PRECURSOR (ATPASE PROTEIN 9)(SUBUNIT C)	
4753	14638	24425	2.71	1.0E-25 BE162737.1	EST_HUMAN	PM1-HT0454-080100-002-109 HT0454 Homo sapiens cDNA	
6089	19460	26199	2.85	1.0E-25 AA5822690.1	EST_HUMAN	nn54h11.s1 Scars_fetal_heart_NbHH19W Homo sapiens cDNA clone MAGE:1087749 3'	
6616	16496	26983	3.15	1.0E-25 AA709079.1	EST_HUMAN	Zf9cg04.s1 Scars_fetal_heart_NbHH19W Homo sapiens cDNA clone MAGE:384822 3' similar to contains PTR5.13 PTR5 repetitive element;	
8337	18214	28467	3.5	1.0E-25 U93163.1	NT	Homo sapiens MAGE-B2 (MAGE-B2), MAGE-B3 (MAGE-B3), MAGE-B4 (MAGE-B4), and MAGE-B1 (MAGE-B1) genes, complete cds	
9143	18897	28796	1.45	1.0E-25 D14547.1	NT	Human DNA, SINE repetitive element	
9143	18897	28797	1.45	1.0E-25 D14547.1	NT	Human DNA, SINE repetitive element	
9914	19389		1.32	1.0E-25 X51755.1	NT	Human lambda-immunglobulin constant region complex (germline)	
2433	12310	22206	1.94	9.0E-26 AL163218.2	NT	Human sapiens chromosome 21 segment HS21C018	
9010	19501		1.73	9.0E-26 AL163285.2	NT	Human sapiens chromosome 21 segment HS21C035	
5492	15411		1.56	8.0E-26 D14547.1	NT	Human DNA, SINE repetitive element;	
1559	11464	21321	1.44	7.0E-26 AF003528.1	NT	Homo sapiens X-linked arthriotic ectodermal dysplasia protein gene (EDA), exon 2 and flanking repeat regions	
3898	13808	23694	1.35	7.0E-26 X89211.1	NT	H. sapiens DNA for endogenous retroviral like element	
4067	13969	23745	2.03	7.0E-26 AW340153.1	EST_HUMAN	hd02e12.x1 Scars_NFL_T_GBC_S1 Homo sapiens cDNA clone MAGE:2908368 3'	
8923	18731		7.99	7.0E-26 AA115895.1	EST_HUMAN	zn3008.r1 Stratagene neuroepithelium NT2RAMI 937234 Homo sapiens cDNA clone MAGE:5418943 5'	
9700	19250		1.33	7.0E-26 AW954559.1	EST_HUMAN	EST366629 MAGE sequences, MAGC Homo sapiens cDNA	
2178	12065	21967	2.44	6.0E-26 AF029308.1	NT	Human sapiens chromosome 9 duplication of the T cell receptor beta locus and trypsinogen gene families	
3302	13223	23025	1.42	6.0E-26 AA206131.1	EST_HUMAN	zg52h04.r1 Stratagene neuroepithelium #937231 Homo sapiens cDNA clone IMAGE:645271 5'	
8932	18740	23033	4.98	6.0E-26 AL163210.2	NT	Human sapiens chromosome 21 segment HS21C010	

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Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
1160 11073	20918	3.33	5.0E-26	AI708235.1	EST_HUMAN	as38h08_x1 Barstead aorta HPLRB6 Homo sapiens cDNA clone IMAGE:2319519 3' similar to WP:49C12.11 CE03371;	
1160 11073	20919	3.33	5.0E-26	AI708235.1	EST_HUMAN	as38h08_x1 Barstead aorta HPLRB6 Homo sapiens cDNA clone IMAGE:2319519 3' similar to WP:49C12.11 CE03371;	
9976 19435		1.71	5.0E-26	AI761429.1	EST_HUMAN	wg65edc1 Scores_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE:2339986 3' similar to contains Alu repetitive element;	
1525 11430		1.52	4.0E-26	AA329548.1	EST_HUMAN	EST33446 Embryo, 12 week II Homo sapiens cDNA 5' end	
7416 17283		3.77	4.0E-26	7657670 NT	Homo sapiens upstream binding transcription factor, RNA polymerase I (UBTF), mRNA	Homo sapiens upstream binding transcription factor, RNA polymerase I (UBTF), mRNA	
8053 17944	28194	3.74	4.0E-26	BE266187.1	EST_HUMAN	601191345F1 NIH_MGC_7 Homo sapiens cDNA clone IMAGE:3535210 5'	
1958 11853	21740	1.5	3.0E-26	AI045855.2	EST_HUMAN	DKFZp434J066_r1_434 (synonym: hte3) Homo sapiens cDNA clone DKFZp434J066 5'	
1987 11880		2.41	3.0E-26	AA115895.1	EST_HUMAN	zn30c08_r1 Strategene neuroepithelium NT2EM1 937234 Homo sapiens cDNA clone IMAGE:548943 5' similar to qb:M14338 VITAMIN K-DEPENDENT PROTEIN SPRECURSOR (HUMAN);	
3712 13625	23408	1.19	3.0E-26	AA152464.1	EST_HUMAN	z03010_r1 Strategene colon (#837204) Homo sapiens cDNA clone IMAGE:588427 5' similar to TR:G695374	
3712 13625	23409	1.19	3.0E-26	AA152464.1	EST_HUMAN	G695314 THYROID RECEPTOR INTERACTOR;	
3712 13625	23409	1.19	3.0E-26	BF245458.1	EST_HUMAN	z03010_r1 Strategene colon (#837204) Homo sapiens cDNA clone IMAGE:588427 5' similar to TR:G695374	
6131 15978	26114	4.35	3.0E-26	AF036405.1	NT	G695314 THYROID RECEPTOR INTERACTOR;	
8108 17998		2.18	3.0E-26	AF036405.1	(Homo sapiens MLL (MLL) gene, exons 1-3, and partial cds)	601854963F1 NIH_MGC_57 Homo sapiens cDNA clone IMAGE:4083278 5'	
8853 18665	28951	1.99	3.0E-26	AW875681.1	EST_HUMAN	QV2-PT0012-040400-124-e05 PT0012 Homo sapiens cDNA	
8853 18665	28952	1.99	3.0E-26	AW875681.1	EST_HUMAN	QV2-PT0012-040400-124-e05 PT0012 Homo sapiens cDNA	
8879 18691	28984	10.55	3.0E-26	AA583173.1	EST_HUMAN	nn37d05_s1 NCL_CGAP_GC5 Homo sapiens cDNA clone IMAGE:1088057 3' similar to contains OFR:t1 OFR repetitive element;	
666 10600	20418	5.61	2.0E-26	AI163282.2	NT	Homo sapiens chromosome 21 segment HS21CC082	
1825 11722		3.36	2.0E-26	AI038099.2	EST_HUMAN	DKFZp568L171_s1_566 (synonym: hkd2) Homo sapiens cDNA clone DKFZp568L171 3'	
3193 13118	22924	4.26	2.0E-26	X86694.1	NT	M.musculus mRNA for astrocytic phosphoprotein, PEA-15	
8135 18023		2.88	2.0E-26	D87675.1	NT	Homo sapiens DNA for amyloid precursor protein, complete cds	
8549 18420	28690	4.55	2.0E-26	AI801412.1	EST_HUMAN	t089a01_x1 NC1_CGAP_Gas4 Homo sapiens cDNA clone IMAGE:2185446 3' similar to contains Alu repetitive element; contains element MER20 MER20 repetitive element;	
8720 18537		1.82	2.0E-26	AF055096.1	NT	Homo sapiens MHC class 1 region	
9252 18962		2.19	2.0E-26	AB378859.1	NT	Homo sapiens mRNA for KIAA1438 protein, partial cds	
132 10106	19927	2.36	1.0E-26	BE170371.1	EST_HUMAN	QV4-HT0538-020300-123-a02 HT0538 Homo sapiens cDNA	
2001 11894	21786	1.39	1.0E-26	AI039363.2	EST_HUMAN	DKFZp34H910_r1_434 (synonym: hte3) Homo sapiens cDNA clone DKFZp34H910 5'	
2516 12390	22282	0.84	1.0E-26	BE814995.1	EST_HUMAN	MR2-BN0114-240500-030-g07 BN0114 Homo sapiens cDNA	
2654 12521		16.79	1.0E-26	AF261085.1	NT	Homo sapiens glyceraldehyde-3-phosphate dehydrogenase (GADPH) mRNA, complete cds	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
6089	16034		2.75	1.0E-26 BE 65980.1	EST_HUMAN	MR3-HT0487-150200-113-g01 HT0487 Homo sapiens cDNA	
8266	18146		3.17	1.0E-26 AL038487.1	EST_HUMAN	DKFZp566C2146_1' 566 (synonym: hifid2) Homo sapiens cDNA clone DKFZp566C2146 5'	
9483	19737		1.84	1.0E-26 HS60893.1	EST_HUMAN	CHR220032 Chromosome 22 exon Homo sapiens cDNA clone C22_45 5'	
7364	17342		3.11	9.0E-27 U93163.1	NT	Homo sapiens MAGE-B2 (MAGE-B2), MAGE-B3 (MAGE-B3), MAGE-B4 (MAGE-B4), and MAGE-B1 (MAGE-B1) genes, complete cds	
8027	19470	28118	3.48	9.0E-27 P54296	SWISSPROT	MYOMESIN 2 (M-PROTEIN) (165 KD TITIN-ASSOCIATED PROTEIN) (165 KD CONNECTIN-ASSOCIATED PROTEIN)	
9013	18812		3.97	9.0E-27 BE45556.1	EST_HUMAN	na03c07.x1 NCL_CGAP_Pr23 Homo sapiens cDNA clone IMAGE:3253644 3' similar to contains OFR.t1 OFR repetitive element;	
10	9996	19787	3.09	8.0E-27 AI831462.1	EST_HUMAN	wI49c04.x1 NCL_CGAP_Lu19 Homo sapiens cDNA clone IMAGE:2406150 3' similar to contains THR.b2 THR repetitive element;	
545	10486		4.13	8.0E-27 AL163227.2	NT	Homo sapiens chromosomes 21 segment HS21C027	
1395	11300	21158	18.87	8.0E-27 AW162737.1	EST_HUMAN	au87h08.x1 Schneider fetal brain 00004 Homo sapiens cDNA clone IMAGE:2783295 3' similar to gb:K00558 TUBULIN ALPHA-1 CHAIN (HUMAN);	
1395	11300	21159	18.87	8.0E-27 AW162737.1	EST_HUMAN	au87h08.x1 Schneider fetal brain 00004 Homo sapiens cDNA clone IMAGE:2783295 3' similar to gb:K00558 TUBULIN ALPHA-1 CHAIN (HUMAN);	
2121	12009	21909	0.98	8.0E-27 AW834776.1	EST_HUMAN	PM2-SN0018-220500-002-a07 SN0018 Homo sapiens cDNA ADP,ATP CARRIER PROTEIN, LIVER ISOFORM T2 (ADP/ATP TRANSLOCASE 3) (ADENINE NUCLEOTIDE TRANSLOCATOR 3) (ANT 3)	
3148	13073	22874	3.31	8.0E-27 P12236	SWISSPROT	NUCLEOTIDE TRANSLOCATOR 3 (ANT 3)	
3309	13230	23035	0.91	8.0E-27 AF181897.1	NT	Homo sapiens WRN (WRN) gene, complete cds	
6165	15122		3.12	8.0E-27 BE926560.1	EST_HUMAN	MR4-BT0398-250800-204-d06 BT0398 Homo sapiens cDNA	
6196	15956	26088	4	8.0E-27 N84970.1	EST_HUMAN	J1751F Human fetal heart, Lambda ZAP Express Homo sapiens cDNA clone J1751 5' similar to REPETITIVE ELEMENT L1	
7310	17186	27286	1.68	8.0E-27 AW837579.1	EST_HUMAN	CM1-CT0315-091299-063-d07 CT0315 Homo sapiens cDNA	
7310	17186	27387	1.68	8.0E-27 AW837579.1	EST_HUMAN	CM1-CT0315-091299-063-d07 CT0315 Homo sapiens cDNA	
668	10602		1.22	7.0E-27 Z70664.1	NT	Human endogenous retroviral element HC2	
5023	14896		2.09	7.0E-27 AW629172.1	EST_HUMAN	hi51h12.x1 Seares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2975879 3' similar to TR:076040 O76040 ORF2, FUNCTION UNKNOWN; ;	
8132	18020		4.22	7.0E-27 AJ271735.1	NT	Homo sapiens Xq pseudoautosomal region, segment 1/2	
9651	19204		2.07	7.0E-27 AV723365.1	EST_HUMAN	AV723365 HTB Homo sapiens cDNA clone HTBAHE02 5'	
8109	17999	28246	6.21	6.0E-27 M26697.1	NT	Human nucleolar protein (B23) mRNA, complete cds	
7877	17727	27910	2.92	5.0E-27 BF666614.1	EST_HUMAN	60212491F1 NIH MGCG 56 Homo sapiens cDNA clone IMAGE:4278527 5'	
7877	17727	27971	2.92	5.0E-27 BF666614.1	EST_HUMAN	60212491F1 NIH MGCG 56 Homo sapiens cDNA clone IMAGE:4278527 5'	
6046	15949	26080	1.54	4.0E-27	9910569 NT	Mus musculus sperm tail associated protein (Sap), mRNA	

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
6656	16536			1.23	4.0E-27 AF078779.1	NT	Rattus norvegicus putative four repetition channel mRNA, complete cds
8880	18632	28985	2.56	4.0E-27 X89211.1	NT	H_sapiens DNA for endogenous retroviral like element	
1995	11889	21782	5.42	3.0E-27 X60658.1	NT	R_rattus RY43 mRNA for a potential ligand-binding protein	
4174	14074	23849	1.27	3.0E-27 BE071924.1	EST_HUMAN	PM0-BT0527-090100-001-d11 BT0527_Homo sapiens cDNA	
5278	15200	24976	5.13	3.0E-27 AA077705.1	EST_HUMAN	7B44C08 Chromosome 7 Fetal Brain cDNA Library Homo sapiens cDNA clone 7B44C08	
7365	17343	27549	2.98	3.0E-27 BT035327.1	EST_HUMAN	601458531FI_NIH_MGC_66 Homo sapiens cDNA clone IMAGE:3862086 5'	
36	10023	19820	7.96	2.0E-27 AF054187.1	NT	Homo sapiens alpha NAC mRNA, complete cds	
1853	11749		18.58	2.0E-27 AA1665345.1	EST_HUMAN	nk01b10.s1 NCI_CGAP_Pri1 Homo sapiens cDNA clone IMAGE:1000699 similar to gb:M17886 60S	
3071	12998		10.27	2.0E-27 AW629172.1	EST_HUMAN	h51lh12.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2975879 3' similar to TR:O76040	
3187	13112	22916	1.43	2.0E-27 AF111167.2	NT	Homo sapiens jun dimerization protein gene, partial cds; cfos gene, complete cds; and unknown gene	
3187	13112	22917	1.43	2.0E-27 AF111167.2	NT	Homo sapiens jun dimerization protein gene, partial cds; cfos gene, complete cds; and unknown gene	
3944	13852	23627	1.09	2.0E-27 AF00368.1	NT	Rattus norvegicus voltage-gated sodium channel mRNA, complete cds	
6712	16592	26780	1.5	2.0E-27 AI866347.1	EST_HUMAN	W128g07.x1 NCI_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2426268 3'	
7338	17206		2.3	2.0E-27 AA5151527.1	EST_HUMAN	nk08h05.s1 NCI_CGAP_Thy1 Homo sapiens cDNA clone IMAGE:943737 similar to contains L1.k3 L1 repetitive element;	
7768	17618	27847	1.22	2.0E-27 M78590.1	EST_HUMAN	EST070736 Fetal brain, Strategene (cat#936206) Homo sapiens cDNA clone HFBCF07	
7768	17618	27848	1.22	2.0E-27 M78590.1	EST_HUMAN	EST070738 Fetal brain, Strategene (cat#936206) Homo sapiens cDNA clone HFBCF07	
8324	18201	28450	2.89	2.0E-27 AU121685.1	EST_HUMAN	AUJ21655 Mammal Homo sapiens cDNA clone MMAMA1000746 5'	
8778	11749		20.82	2.0E-27 AA1665345.1	EST_HUMAN	nk01b10.s1 NCI_CGAP_Pri1 Homo sapiens cDNA clone IMAGE:1000699 similar to gb:M17886 60S	
429	10374		1.56	1.0E-27 AL103246.2	NT	Homo sapiens chromosome 21 segment HS21C046	
980	10903	20749	1.41	1.0E-27 AB026898.1	NT	Homo sapiens DNA, DLECT1 to ORC1L4 gene region, section 1/2 (DLECT1, ORC1L3, ORC1L4 genes, complete cds)	
5943	15848	25972	6.31	1.0E-27 8005885 NT	NT	Homo sapiens Retina-derived POU-domain factor-1 (RPF-1), mRNA	
6105	15999	26136	2.2	1.0E-27 F30158.1	EST_HUMAN	HSPD20461 HM3 Homo sapiens cDNA clone s4000095C10	
6105	15999	26137	2.2	1.0E-27 F30158.1	EST_HUMAN	HSPD20461 HM3 Homo sapiens cDNA clone s4000095C10	
7005	16882	27075	1.72	1.0E-27 AB007923.1	NT	Homo sapiens mRNA for KIAA0454 protein, partial cds	
7194	17071		1.89	1.0E-27 BE079780.1	EST_HUMAN	RCG-BT0627-140200-011-E06 BT0627 Homo sapiens cDNA	
7606	17457	27672	2.68	1.0E-27 D874749.1	NT	Human mRNA for KIAA0260 gene, partial cds	
8943	18751	29046	3.14	1.0E-27 AF111093.1	NT	Bos taurus latrophilin 3 splice variant bbab mRNA, complete cds	

Table 4
Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
136	10109		2.32	9.0E-28	BE248399.1	EST_HUMAN	hw17c1.x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3188188 3' similar to TR:Q07314 Q07314
303	10270	20089	3.01	9.0E-28	AU126260	EST_HUMAN	SECRETED NEUREXIN III-ALPHA-C PRECURSOR. [3] TR:Q07280 TR:Q07313 ;
5118	14986	24760	1.21	9.0E-28	AI590115.1	EST_HUMAN	AU126260 NT2RP1 Homo sapiens cDNA clone NT2RP1000443 5'
5118	14986	24761	1.21	9.0E-28	AI590115.1	EST_HUMAN	to12609.x1 NCI_CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2178809 3' similar to contains OFR.t1 OFR repetitive element;
9093	18886		3.74	9.0E-28	BF377859.1	EST_HUMAN	to12609.x1 NCI_CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2178809 3' similar to contains OFR.t1 OFR repetitive element;
9418	19592		1.97	8.0E-28	AW157571.1	EST_HUMAN	CM2-TN0140-070900-372.g01 TN0140 Homo sapiens cDNA
1164	11077	20922	7.54	7.0E-28	AU142750.1	EST_HUMAN	au83h05.x1 Schneider fetal brain 0004 Homo sapiens cDNA clone IMAGE:2782911 3' similar to TR:O60302 KIAA0555 PROTEIN; contains element MER22 repetitive element;
8523	18395	28660	2.54	7.0E-28	11417865	NT	AU142750 Y79AA1 Homo sapiens cDNA clone YT9AA1000824 5'
9053	18336		1.44	7.0E-28	AV735348.1	EST_HUMAN	Homo sapiens gamma-glutamyltranspeptidase-like activity 1 (GGT1A1), mRNA
3987	13594	23670	1.27	6.0E-28	AB020673.1	NT	AV735348 CB Homo sapiens cDNA clone CBFAKA12 5'
3987	13594	23671	1.27	6.0E-28	AB020673.1	NT	Homo sapiens mRNA for KIAA0866 protein, complete cds
9673	19232		2.73	6.0E-28	AA504562.1	EST_HUMAN	aad0d03.r1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:825340 5' similar to contains Alu repetitive element; contains element PTR5 repetitive element;
315	10277		3.08	5.0E-28	AI921003.1	EST_HUMAN	w018c07.x1 NCI_CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2455692 3' similar to contains THR.b1 THR repetitive element;
3927	138336	23616	1.85	5.0E-28	R79762.1	EST_HUMAN	yf8f10.r1 Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE:146443 5'
2584	12455	22347	1.42	4.0E-28	AW195066.1	EST_HUMAN	xr33c09.x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2685504 3' similar to SW:GG95_HUMAN Q053379 GOLGIN-95 ;
3070	12997	22788	3.39	4.0E-28	BE409100.1	EST_HUMAN	601300703F NIH MGCC_21 Homo sapiens cDNA clone IMAGE:36335305 5'
6336	16199	26359	1.59	4.0E-28	AI198941.1	EST_HUMAN	qf68f10.x1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:1755019 3' similar to gb:M19503 LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (HUMAN);
8239	18119		4.29	4.0E-28	AF029308.1	NT	Homo sapiens chromosome 9 duplication of the T cell receptor beta locus and trypsinogen gene families
8371	18248		53.6	4.0E-28	AB035241.1	NT	Felis catus GAPDH mRNA for glyceraldehyde-3-phosphate dehydrogenase, complete cds
8388	16199	26359	2.94	4.0E-28	AI198941.1	EST_HUMAN	qf68f10.x1 Soares testis_NHT Homo sapiens cDNA clone IMAGE:1755019 3' similar to gb:M19503 LINE-1 REVERSE TRANSCRIPTASE HOMOLOG (HUMAN);
1262	11169		1.89	3.0E-28	AF155382.1	NT	Homo sapiens metallopeptidase-like, disintegrin-like, cysteine-rich protein 2 epsilon (ADAM22) mRNA, complete cds
7124	17001	27193	2.19	3.0E-28	BF354030.1	EST_HUMAN	MFR3-HT013-280500-013-09 HT0713 Homo sapiens cDNA
8306	18183	28430	1.84	3.0E-28	U53388.1	NT	Homo sapiens MHC class 1 region

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
9489	19106		2.44	3.0E-28	AI831991.1	EST_HUMAN	wj98107_x1 NCI_CGAP_Lym12 Homo sapiens cDNA clone IMAGE:2410885 3' similar to contains Alu repetitive element; contains element HGR repetitive element;
83	10067	19884	6.45	2.0E-28	BE062167.1	EST_HUMAN	RC1-BT0254-220300-019-cd5 BT0254 Homo sapiens cDNA
1149	11062	20905	10	2.0E-28	Y11107.3	NT	Homo sapiens ITGB4 gene for integrin beta 4 subunit, exons 3-41
2427	12304	22200	2.1	2.0E-28	AI318634.1	EST_HUMAN	qo35b03_x1 NCI_CGAP_Lu6 Homo sapiens cDNA clone IMAGE:1910483 3' similar to contains L1,b2L1 repetitive element;
5821	15727		4.28	2.0E-28	BF7212905.1	EST_HUMAN	601814196F1 NIH_MGC_54 Homo sapiens cDNA clone IMAGE:4048751 5'
7528	17379		5.86	2.0E-28	AW972305.1	EST_HUMAN	EST3843394 IMAGE sequences, MAGL Homo sapiens cDNA
8887	18698	28992	2.27	2.0E-28	AF224669.1	NT	Homo sapiens mannosidase, beta A, lysosomal (MANBA) gene, and ubiquitin-conjugating enzyme E2D 3 (UBE2D3) genes, complete cds
1483	11368	21233	2.42	1.0E-28	D38044.1	NT	Human gene for Ah-receptor, exon 7-9
2173	12060	21953	1.55	1.0E-28	BF3133236.1	EST_HUMAN	QV1-BT0821-120900-360-b03 BT0821 Homo sapiens cDNA
2650	12517	22407	1.03	1.0E-28	AF000995.1	NT	Homo sapiens ubiquitous TPR motif, Y isoform (UTY) mRNA, alternative transcript 2, complete cds
6587	16467		4.48	1.0E-28	11429835	NT	Homo sapiens similar to ribosomal protein L12 (H. sapiens) (LOC63091), mRNA
6675	16555		3.02	1.0E-28	8922793	NT	Homo sapiens hypothetical protein FLJ10968 (FLJ10968), mRNA
7346	17214	27413	2.63	1.0E-28	AA303744.1	EST_HUMAN	EST179615 HCC cell line (metastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to retroviral LTR
7678	17528	27753	5.15	1.0E-28	4758431	NT	Homo sapiens gamma-glutamyltransferase-like activity 1 (GGTLA1), mRNA
7678	17528	27754	5.15	1.0E-28	4758431	NT	Homo sapiens gamma-glutamyltransferase-like activity 1 (GGTLA1), mRNA
9058	18840		3.95	1.0E-28	AA054182.1	EST_HUMAN	Zf51071.1 Soares retina N2b4-HR Homo sapiens cDNA clone IMAGE:3804418 5'
9797	19485		1.43	1.0E-28	AL153247.2	NT	Homo sapiens chromosome 21 segment HS21C047
9892	19691	24896	2.6	9.0E-29	AW663987.1	EST_HUMAN	hi76906_x1 Soares NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:297B266 3'
9585	19169		2.92	8.0E-29	Q00130	SWISSPROT	HYPOTHETICAL GENE 50 PROTEIN
1584	11488	21349	0.91	7.0E-29	AW366447.1	EST_HUMAN	EST37821 IMAGE sequences, MAGI Homo sapiens cDNA
9942	19410		5.08	7.0E-29	AJ123352.1	NT	Rattus norvegicus mRNA for 45 kDa secretory protein, partial
580	10518	20325	6.52	6.0E-29	AI936748.1	EST_HUMAN	Q15475 UNNAMED HERV-H PROTEIN ;contains LTR7_b1 LTR7 repetitive element;
9353	19028		3.88	6.0E-29	BE940436.1	EST_HUMAN	RC3-UT0062-210800-021-c05 UT0062 Homo sapiens cDNA
9438	19076		1.86	6.0E-29	BF568097.1	EST_HUMAN	602184092F1 NIH_MGC_42 Homo sapiens cDNA clone IMAGE:43000079 5'
4928	14807		0.98	5.0E-29	AL163203.2	NT	Homo sapiens chromosome 21 segment HS21C03
7068	16945		7.61	5.0E-29	AW887541.1	EST_HUMAN	RC3-OT0091-170300-011-c12 OT0091 Homo sapiens cDNA
3194	13119		1.68	4.0E-29	AI752367.1	EST_HUMAN	cn15c02_x1 Normal Human Trabecular Bone Cells Homo sapiens cDNA clone NHTBC_cn15c02 random
5656	15568		6.65	4.0E-29	BE164930.1	EST_HUMAN	QV1-HT0471-280300-121-a05 HT0471 Homo sapiens cDNA

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
7079	16956	27149	4.71	4.0E-29	J04988.1	NT	Human 90 kD heat shock protein gene, complete cds
4317	14244	23997	1.45	3.0E-29	AB042297.1	NT	Homo sapiens PTS gene for 6-pyruvoyltetrahydropterin synthase, complete cds
4626	14514	24305	1.31	3.0E-29	BF533236.1	EST_HUMAN	QV1-BT0821-1/20300-360-b03 BT0821 Homo sapiens cDNA
7070	16947	27138	2.07	3.0E-29	D38044.1	NT	Human gene for Ah-receptor, exon 7-9
7362	17229	27429	1.6	3.0E-29	AW303817.1	EST_HUMAN	x17703_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2813405 3' similar to contains Ali repetitive element; contains MER19/12 MER19 repetitive element;
7492	17362		1.64	3.0E-29	AL163246.2	NT	Homo sapiens chromosome 21 segment HS21C046
8591	18459	28728	2.22	3.0E-29	AA0403033.1	EST_HUMAN	z62901_x1 Soares_festis_NHT Homo sapiens cDNA clone IMAGE:726889 5' similar to TR:G1335769 G1335769 GAC-POL POLYPROTEIN ;
9248	18959		1.51	3.0E-29	D63882.1	NT	Human Hs1M15 mRNA for Hs1M15, complete cds
484	10427	20240	1.43	2.0E-29	AF084869.1	NT	Homo sapiens envelope protein RIC-6 (env) gene, complete cds
484	10427	20241	1.43	2.0E-29	AF084869.1	NT	Homo sapiens envelope protein RIC-6 (env) gene, complete cds
1516	11421	21278	6.12	2.0E-29	AI963604.1	EST_HUMAN	wf65d10_x1 NCL_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2492563 3' similar to TR:O16546 O15546 HERV-E ENVELOPE GLYCOPROTEIN ;
1516	11421	21279	6.12	2.0E-29	AI963604.1	EST_HUMAN	wf65d10_x1 NCL_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2492563 3' similar to TR:O16546 O15546 HERV-E ENVELOPE GLYCOPROTEIN ;
4182	14082	23885	2.03	2.0E-29	AL163268.2	NT	Homo sapiens chromosome 21 segment HS21C068
5750	15658	25765	1.43	2.0E-29	AI806418.1	EST_HUMAN	wf27g07_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2356860 3' similar to contains element MER6 repetitive element;
6461	15658	25765	1.43	2.0E-29	AI806418.1	EST_HUMAN	wf27g07_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2356860 3' similar to contains element MER6 repetitive element;
7474	17334	27539	2.95	2.0E-29	AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048
7474	17334	27540	2.95	2.0E-29	AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048
7879	17729	27973	3.39	2.0E-29	AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048
7879	17729	27974	3.39	2.0E-29	AL163248.2	NT	Homo sapiens chromosome 21 segment HS21C048
8804	18618		1.96	2.0E-29	AW380701.1	EST_HUMAN	QV0-OT0032-080300-155-d01 OT0032 Homo sapiens cDNA
9004	18807		1.73	2.0E-29	AL163227.2	NT	Homo sapiens chromosome 21 segment HS21C027
7105	16982	27174	5.41	1.0E-29	AW383880.1	EST_HUMAN	RC1-HN0003-220300-021-b04 HN0003 Homo sapiens cDNA
5966	15871	25995	3.04	9.0E-30	AA761215.1	EST_HUMAN	n220c07_s1 NCL_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:1288332 3' similar to contains MER4b1 MER4 repetitive element;
9132	18889		1.5	9.0E-30	11422745	NT	Homo sapiens zinc/furan regulated transporter-like (ZIRTL), mRNA
5815	15721		8.81	8.0E-30	FO8888.1	EST_HUMAN	HSC23F051 normalized infant brain cDNA Homo sapiens cDNA clone c-23f051
6812	16691	26880	3.37	8.0E-30	AA383873.1	EST_HUMAN	EST79737 Thymus I Homo sapiens cDNA 5' end similar to EST containing O family repeat
7039	16916	27105	3.53	8.0E-30	AI657072.1	EST_HUMAN	PT2.1_13_B11r tumor2 Homo sapiens cDNA 3'

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit No.	Top Hit Database Source	Top Hit Descriptor
1499	11403		1.16	7.0E-30	BE091133.1	EST_HUMAN	PM4-BT0724-150400-004-d11 BT0724 Homo sapiens cDNA
1549	11454		0.95	6.0E-30	X51755.1	NT	Human lambda-immunoglobulin constant region complex (germline)
1736	11637	21505	1.26	6.0E-30	D25303.1	NT	Human mRNA for integrin alpha subunit, complete cds
3153	13078	22878	2.41	6.0E-30	BE008026.1	EST_HUMAN	QV0-BN0147-290400-214-f12 BN0147 Homo sapiens cDNA
9915	11454		3.15	6.0E-30	X51755.1	NT	Human lambda-immunoglobulin constant region complex (germline)
3931	13840	23520	31.23	5.0E-30	A1399892.1	EST_HUMAN	tg2903.x1 NCL CGAP_CLL1 Homo sapiens cDNA clone IMAGE:21162763 similar to contains Alu repetitive element
5205	19522		7.35	5.0E-30	U87981.1	NT	Human aconitase hydratase (ACO2) gene, exon 7
8261	18141		3.55	5.0E-30	AL163278.2	NT	Human sapiens chromosome 21 segment HS21C078
8495	18368	28632	6.68	5.0E-30	AL163210.2	NT	Human sapiens chromosome 21 segment HS21C010
8495	18368	28633	6.68	5.0E-30	AL163210.2	NT	Human sapiens chromosome 21 segment HS21C010
2096	11985	21880	1.64	4.0E-30	AV937471.1	EST_HUMAN	Q173-DT0043-090200-050-c06 DT0043 Homo sapiens cDNA
2096	11985	21881	1.64	4.0E-30	AV937471.1	EST_HUMAN	QV3-DT0043-090200-050-c06 DT0043 Homo sapiens cDNA
7162	17039	27231	1.49	4.0E-30	AV812488.1	EST_HUMAN	CM1-ST0181-091199-055-ST0181 Homo sapiens cDNA
							qq93c5-x1 Seares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:19389203' similar to contains MER29,b2 MER29 repetitive element;
1134	11048		1.71	3.0E-30	A1338551.1	EST_HUMAN	
3697	13611	23395	0.91	3.0E-30	AF128893.1	NT	Hom sapiens telomerase reverse transcriptase (TERT) gene, exons 1-6
8538	18410	28675	2.38	3.0E-30	F34056	SWISSPROT	TRANSCRIPTION FACTOR AP-2
659	10593	20411	1.18	2.0E-30	AV857315.1	EST_HUMAN	CM0-CT0307-310100-158-h03 CT0307 Homo sapiens cDNA
1068	10984		2.32	2.0E-30	F08688.1	EST_HUMAN	HSC23F051 normalized infant brain cDNA Homo sapiens cDNA clone c-23f05
1464	11369	21234	5.91	2.0E-30	BE175877.1	EST_HUMAN	RC5-HT0582-1110400-013-H08 HT0582 Homo sapiens cDNA
2683	12548	22438	10.97	2.0E-30	BE768232.1	EST_HUMAN	IL2-NT0101-280700-116-EG4 NT0101 Homo sapiens cDNA
2889	12816	22609	6.38	2.0E-30	AF114156.1	NT	Hom sapiens Y-linked zinc finger protein (ZFY) gene, complete cds
3721	13633	23419	2.11	2.0E-30	AV206561.1	EST_HUMAN	UH-B11-efo-c-12-0-U1-s1 NG1_CGAP_Sub3 Homo sapiens cDNA clone IMAGE:27225533
4668	14554	24346	1.72	2.0E-30	BE298945.1	EST_HUMAN	6011198860F1 NIH_MGC_17 Homo sapiens cDNA clone IMAGE:30294385
4668	14554	24347	1.72	2.0E-30	BE298945.1	EST_HUMAN	6011198860F1 NIH_MGC_17 Homo sapiens cDNA clone IMAGE:30294385
6966	16844	27036	3.46	2.0E-30	C18939.1	EST_HUMAN	C18939 Human placenta cDNA (T-Fujiiwara) Homo sapiens cDNA clone GENL-570C01 5'
7019	16896	27085	1.55	2.0E-30	BE670617.1	EST_HUMAN	T637c12.x1 NCL CGAP_Lu24 Homo sapiens cDNA clone IMAGE:32846623 similar to SW:DHS_A HUMAN P31040 SUCCINATE DEHYDROGENASE [UBIQUINONE] FLAVOPROTEIN SUBUNIT PRECURSOR;
7019	16896	27086	1.55	2.0E-30	BE670617.1	EST_HUMAN	T637c12.x1 NCL CGAP_Lu24 Homo sapiens cDNA clone IMAGE:32846623 similar to SW:DHS_A HUMAN P31040 SUCCINATE DEHYDROGENASE [UBIQUINONE] FLAVOPROTEIN SUBUNIT PRECURSOR;
7743	17593	27814	3.3	2.0E-30	AW971558.1	EST_HUMAN	EST383657 MAGE resequences, MAGL Homo sapiens cDNA

Table 4

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Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
7802	17652	27889	4.47	2.0E-30	AW470791.1	EST_HUMAN	hs33d06.x1 NCI_CGAP_Kid12 Homo sapiens cDNA clone IMAGE:2875499 3' similar to contains THR:k3
284	10249	20069	10.22	1.0E-30	C18939.1	EST_HUMAN	THR repetitive element; C18939 Human placenta cDNA (T Fujimura) Homo sapiens cDNA clone GEN-5700C01 5'
527	10469	20281	1.6	1.0E-30	AW468897.1	EST_HUMAN	hd30b04.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2910891 3' similar to contains MER1:k3 MER1 repetitive element;
699	10632	20457	2.92	1.0E-30	AL163203.2	NT	Homo sapiens chromosome 21 segment HS21C003
2165	12052	21953	3.24	1.0E-30	AA664377.1	EST_HUMAN	ac77b08.s1 Strategene lung (#937210) Homo sapiens cDNA clone IMAGE:858599 3'
2413	12280	22187	2.57	1.0E-30	BF547728.1	EST_HUMAN	60202260F1 NCI_CGAP_Bm67 Homo sapiens cDNA clone IMAGE:4157901 5'
2973	12300	22699	1.03	1.0E-30	58030391	NT	Homo sapiens methionine aminopeptidase; eIF-2-associated p67 (MNPFP), mRNA
3015	12943	22736	0.93	1.0E-30	AA315045.1	EST_HUMAN	EST186868 HCC cell line (metastasis to liver in mouse) II Homo sapiens cDNA 5' end
6545	16403	26582	2.32	1.0E-30	BF183230.1	EST_HUMAN	60180932F1 NIH_MGC_18 Homo sapiens cDNA clone IMAGE:404694 5'
9733	19615		5.1	1.0E-30	HS5593.1	EST_HUMAN	CHR220532 Chromosome 22 exon Homo sapiens cDNA clone C22_728 5'
7013	16890		2.04	9.0E-31	Z38298.1	EST_HUMAN	HSC05f032 normalized infant brain cDNA Homo sapiens cDNA clone c-05f03 3'
1060	10977	20820	1.64	8.0E-31	8923389	NT	Homo sapiens hypothetical protein FLJ20420 (FL_20420), mRNA
2362	12242		5.14	8.0E-31	AL163208.2	NT	Homo sapiens chromosome 21 segment HS21C008
694	10627		1.72	7.0E-31	AA372637.1	EST_HUMAN	EST84555 Colon adenocarcinoma IV Homo sapiens cDNA 5' end
2633	12500	22392	1.7	7.0E-31	BE226517.1	EST_HUMAN	hw05a11.x1 NCI_CGAP_L124 Homo sapiens cDNA clone IMAGE:3182012 3'
2633	12500	22393	1.7	7.0E-31	BE226517.1	EST_HUMAN	hw05a11.x1 NCI_CGAP_L124 Homo sapiens cDNA clone IMAGE:3182012 3'
9583	19168	25272	2.56	7.0E-31	X51755.1	NT	Human lambda-immunoglobulin constant region complex (germline)
3625	13539		2.79	6.0E-31	AF223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-9, and partial cds, alternatively spliced
6739	16618		3.68	6.0E-31	AF0550066.1	NT	Homo sapiens MHC class 1 region
8120	18008	28255	2.11	6.0E-31	AU119105.1	EST_HUMAN	AU119105 HEMBA1 Homo sapiens cDNA clone HEMBA1005050 5'
9191	18926	25351	1.88	6.0E-31	AW372868.1	EST_HUMAN	RC5-BT0377-031-D12 BT0377 Homo sapiens cDNA
9319	19541		1.78	6.0E-31	BE894488.1	EST_HUMAN	601433087F1 NIH_MGC_72 Homo sapiens cDNA clone IMAGE:3918524 5'
185	10157	19973	2.85	5.0E-31	M60694.1	NT	Homo sapiens type I DNA topoisomerase gene, exon 8
185	10157	19974	2.85	5.0E-31	M60694.1	NT	Homo sapiens type I DNA topoisomerase gene, exon 8
6915	16793		1.21	5.0E-31	BF056540.1	EST_HUMAN	7k06f04.x1 NCI_CGAP_GCG Homo sapiens cDNA clone IMAGE:3443479 3' similar to TRQ13537 Q13537
581	10519		3.78	4.0E-31	A1271735.1	NT	SIMILAR TO POGO ELEMENT; contains L1.1L1 repetitive element;
1594	11408	21357	1.04	4.0E-31	Q10473	SWISSPROT	POLYPEPTIDE N-ACETYLGALACTOSAMINYL TRANSFERASE (PROTEIN-UDP ACETYLGLACTOSEAMINYL TRANSFERASE) (UDP-GALNAc-POLYPEPTIDE; N-
1775	11674		2.59	4.0E-31	AL163280.2	NT	Homo sapiens chromosome 21 segment HS21C080

Table 4
Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
2759	12621			1.38	4.0E-31	5730038B	NT
9604	19186			1.52	4.0E-31	11430273	NT
9718	19259			1.38	4.0E-31	ABD08681.1	NT
2550	12423	22314	1.42	3.0E-31	6005871	NT	Homo sapiens SEC63, endoplasmic reticulum translocon component (<i>S. cerevisiae</i>) like (SEC63L), mRNA
6341	16204	26366	9.03	3.0E-31	4826853	NT	Homo sapiens NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASH1) (NDUFB8) mRNA
6425	16286	26448	1.38	3.0E-31	11420329	NT	Homo sapiens hypothetical protein FLJ10842 (FLJ10842), mRNA
6745	16624		1.95	3.0E-31	AL163206.2	NT	Homo sapiens chromosome 21 segment HS21C006
7525	17376	27585	2.86	3.0E-31	D14523.1	NT	Horse mRNA for ferritin L-chain, complete cds
8028	17920	28167	2.01	3.0E-31	P11174	SWISSPROT	40S RIBOSOMAL PROTEIN S15 (RIG PROTEIN)
8493	18396		6.36	3.0E-31	BF035527.1	EST_HUMAN	601458331F1 NIH_MGC_66 Homo sapiens cDNA clone IMAGE:3862086 5'
1875	11771	21647	1.83	2.0E-31	AW838171.1	EST_HUMAN	QV2-LT0051-260300-111-f03 LT0051 Homo sapiens cDNA
2167	12054	21955	1.31	2.0E-31	AI393388.1	EST_HUMAN	tg44905_x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2111672 3'
2292	12174	22074	2.18	2.0E-31	AL119245.1	EST_HUMAN	DKFZp761G1513_1761 (synonym: hany2) Homo sapiens cDNA clone DKFZp761G1513 5'
2391	12289	22164	4.4	2.0E-31	AA458824.1	EST_HUMAN	aa88ff1_s1 Stratagene fetal retina 937202 Homo sapiens cDNA clone IMAGE:8384413 3 similar to contains MER29.63
5502	15421	25483	3.6	2.0E-31	BE250127.1	EST_HUMAN	THR:t2 THR repetitive element;
							ht09g01_X1_NCI_CGAP_Kidf13 Homo sapiens cDNA clone IMAGE:3146256 3' similar to contains MER29.63
7246	17123		1.8	2.0E-31	AA877764.1	EST_HUMAN	nt06104_s1 NCI CGAP_Co10 Homo sapiens cDNA clone IMAGE:1161055 3' similar to TR:Q13537 Q13537
7309	17185	27385	3.99	2.0E-31	7661535	NT	MER37 TRANSPOSSABLE ELEMENT, COMPLETE CONSENSUS SEQUENCE. ;
7700	17550	27774	1.27	2.0E-31	AV710948.1	EST_HUMAN	Homo sapiens B9 protein (B9), mRNA
7700	17550	27775	1.27	2.0E-31	AV710948.1	EST_HUMAN	AV710948 Clu Homo sapiens cDNA clone CuAA1B07 5'
7797	17647	27883	2.17	2.0E-31	BE408611.1	EST_HUMAN	AV710948 Clu Homo sapiens cDNA clone CuAA1B07 5'
7797	17647	27884	2.17	2.0E-31	BE408611.1	EST_HUMAN	601304125F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:3638310 5'
9292	18983		2	2.0E-31	AF448512.1	NT	601304125F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:3638310 5'
9431	19759		1.81	2.0E-31	AI114527.1	EST_HUMAN	Homo sapiens hexokinase II gene, promoter region
16	10002	19794	8.34	1.0E-31	U03163.1	NT	HA1110 Human fetal liver cDNA library Homo sapiens cDNA
1640	11544	21403	2	1.0E-31	O95371	SWISSPROT	Homo sapiens MAGE-B2 (MAGE-B2), MAGE-B3 (MAGE-B3), MAGE-B4 (MAGE-B4), and MAGE-B1 (MAGE-B1) genes, complete cds
1640	11544	21404	2	1.0E-31	O95371	SWISSPROT	OLFACTOORY RECEPTOR 2C1
1640	11544	21405	2	1.0E-31	O95371	SWISSPROT	OLFACTOORY RECEPTOR 2C1
4535	14428	24209	1.01	1.0E-31	AL134376.1	EST_HUMAN	DKFZp47B235_r1 547 (synonym: hb1r1) Homo sapiens cDNA clone DKFZp47B235 5'

Table 4
Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon ORF SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
4535	14428	24210	1.01	1.0E-31	AL134376.1	EST_HUMAN	DKFZp547B235_1' 547 (synonym: hifb1) Homo sapiens cDNA clone DKFZp547B235 5'
5235	15159	24927	3.15	1.0E-31	AW391679.1	EST_HUMAN	MR3-ST0220-151298-028-a08_1 ST0220 Homo sapiens cDNA
5728	15635	25738	1.87	1.0E-31	AF048727.1	NT	Homo sapiens minisatellite ceb1 repeat region
8288	18167	28411	2.72	1.0E-31	AJ086434.1	EST_HUMAN	qf21h03_x1 NC1 CGAP_Bm25 Homo sapiens cDNA clone IMAGE:1750709 3' similar to TIR:Q16595
2835	12763		0.99	9.0E-32	U59871.1	NT	Human familial Alzheimer's disease (STM2) gene, complete cds
6000	15905	26029	2.34	9.0E-32	AV723976 HTB Homo sapiens cDNA clone HTBAAG01 5'	EST_HUMAN	AV723976 HTB Homo sapiens cDNA clone HTBAAG01 5'
2032	11923	21815	3.32	8.0E-32	AJ056770.1	EST_HUMAN	czf5a09_x1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:1675384 3'
4754	14639	24426	1.17	7.0E-32	P52591	SWISSPROT	NUCLEAR ENVELOPE PORE MEMBRANE PROTEIN POM 121 (PORE MEMBRANE PROTEIN OF 121 KD) (P145)
9268	18972		2.77	7.0E-32	X17283.1	NT	Human chromosome 22 immunoglobulin V(K)1 gene, part. with 5' breakpoint between orphons and neighbouring non-ampelified region
2702	12566	22456	0.9	6.0E-32	AJ478104.1	EST_HUMAN	tm34a10_x1 NC1 CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2159994 3' similar to contains MER29;3
6359	16222		1.5	6.0E-32	BE388016.1	EST_HUMAN	MER29 repetitive element;
1018	10936	20779	29.57	5.0E-32	AF116627.1	NT	601511530F1 NIH_MGC_71 Homo sapiens cDNA clone IMAGE:3913087 5'
9151	10839		1.74	4.0E-32	AL163246.2	NT	Homo sapiens chromosome 21 segment HS2/TC046
6484	16342	26510	2.82	4.0E-32	11432574	NT	Homo sapiens AT-binding transcription factor 1 (ATBF1), mRNA
6484	16342	26511	2.82	4.0E-32	11432574	NT	Homo sapiens AT-binding transcription factor 1 (ATBF1), mRNA
448	10392	20213	3.4	3.0E-32	Y17293.1	NT	Homo sapiens FLI-1 gene, partial
1438	11343	21210	7.67	3.0E-32	AV731500.1	EST_HUMAN	AV731500 HTF Homo sapiens cDNA clone HTFAK027 5'
7404	17271	27477	6.73	3.0E-32	AV758634.1	EST_HUMAN	AV758634 BM Homo sapiens cDNA clone BMFBBH12.5'
7404	17271	27478	6.73	3.0E-32	AV758634.1	EST_HUMAN	AV758634 BM Homo sapiens cDNA clone BMFBBH12.5'
8297	18176	28421	8.08	3.0E-32	AA777621.1	EST_HUMAN	Z95a07_s1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:448500 3' similar to
9294	18995		3.04	3.0E-32	BE279086.1	EST_HUMAN	contains THR_B3 THR repetitive element;
9655	15090	24882	2.43	3.0E-32	5174574	NT	601156285F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:3139701 5'
9655	15090	24883	2.43	3.0E-32	5174574	NT	Homo sapiens myeloid/lymphoid or mixed-lineage leukemia (Trithorax (Drosophila) homolog); translocated to, 4 (MLLT4) mRNA
9802	19314		4.94	3.0E-32	BE279086.1	EST_HUMAN	601156285F1 NIH_MGC_21 Homo sapiens cDNA clone IMAGE:3139701 5'
4783	14667	24454	0.91	2.0E-32	BE286613.1	EST_HUMAN	601173631F1 NIH_MGC_17 Homo sapiens cDNA clone IMAGE:3529159 5'
5900	15806	25929	19.01	2.0E-32	Z38133.1	NT	H.sapiens mRNA for myosin
5900	15806	25930	19.01	2.0E-32	Z38133.1	NT	H.sapiens mRNA for myosin

Table 4
Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
6815	16694	26884	3.41	2.0E-32 AA 114294.1	EST_HUMAN	Zn86c08_r1 Stratagene HeLa cell s3 937216 Homo sapiens cDNA clone IMAGE:5651150 5'	
6815	16694	26885	3.41	2.0E-32 AA 114294.1	EST_HUMAN	Zn86c08_r1 Stratagene HeLa cell s3 937216 Homo sapiens cDNA clone IMAGE:5651150 5'	
9908	19385	25175	1.61	2.0E-32 AV 736449.1	EST_HUMAN	AV736449 CB Homo sapiens cDNA clone CBFBIA08 5'	
9908	19385	25176	1.61	2.0E-32 AV 736449.1	EST_HUMAN	AV736449 CB Homo sapiens cDNA clone CBFBIA08 5'	
3055	12982		1.06	1.0E-32 BE 743299.1	EST_HUMAN	601573027F1 NIH_MGC_9 Homo sapiens cDNA clone IMAGE:3834433 5'	
6198	15953	26090	7.04	1.0E-32	11439789 NT	Homo sapiens chromosome 11 open reading frame 9 (C11ORF9), mRNA	
6997	16874	27065	5.18	1.0E-32 AA 720574.1	EST_HUMAN	hw2192.s1 NCI_CGAP_GCE0 Homo sapiens cDNA clone IMAGE:1241138 3' similar to contains THR:t3 THR repetitive element;	
3435	13352		4.98	9.0E-33 BE 327112.1	EST_HUMAN	hw07c05_x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3182216 3' similar to TR:O88539 O88539 WW DOMAIN BINDING PROTEIN 11.1;	
5867	15773		4.19	9.0E-33 AF 223391.1	NT	Homo sapiens calcium channel alpha1E subunit (CACNA1E) gene, exons 7-49, and partial cds, alternatively spliced	
7103	16980	27172	1.99	9.0E-33 BF 247229.1	EST_HUMAN	602021164F1 NCI_CGAP_Bm67 Homo sapiens cDNA clone IMAGE:4156670 5'	
8177	18065		5.89	9.0E-33 AL 163280.2	NT	Homo sapiens chromosome 21 segment HS21C080	
54	10041	19850	2.06	7.0E-33	503 1736 NT	Homo sapiens short-chain alcohol dehydrogenase family member (HEP27) mRNA	
54	10041	19851	2.06	7.0E-33	503 1736 NT	Homo sapiens short-chain alcohol dehydrogenase family member (HEP27) mRNA	
2115	12004	21902	2.71	7.0E-33 A 590115.1	EST_HUMAN	to12609_x1 NCI_CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2178809 3' similar to contains OFR:t1 OFR repetitive element;	
2612	12480		8.4	7.0E-33 AV 730056.1	EST_HUMAN	AV730056 HTF Homo sapiens cDNA clone HTFAVE06 5'	
2800	11573	21439	1.78	7.0E-33 AV 730015.1	EST_HUMAN	AV730015 HTF Homo sapiens cDNA clone HTFANF08 5'	
3204	13128		12.85	7.0E-33 AW 971307.1	EST_HUMAN	EST383396 MAGE resequences, MAGL Homo sapiens cDNA	
8203	18087	28338	3.9	7.0E-33 BF 247229.1	EST_HUMAN	602021164F1 NCI_CGAP_Bm67 Homo sapiens cDNA clone IMAGE:4156670 5'	
8573	18441	28709	2.15	7.0E-33 AW 971568.1	EST_HUMAN	EST383657 MAGE resequences, MAGL Homo sapiens cDNA	
9274	18978	25322	3.48	7.0E-33 AA 014146.1	EST_HUMAN	nt16b01_s1 NCI_CGAP_Phet Homo sapiens cDNA clone IMAGE:1100881 3' similar to contains L1.t1 L1 repetitive element;	
3676	13590		0.85	6.0E-33 AL 163295.2	NT	Homo sapiens chromosome 21 segment HS21C085	
6988	16865	27059	13.52	6.0E-33 J0 038.1	NT	Human glyceraldehyde-3-phosphate dehydrogenase (GAPDH) gene, complete cds	
7049	16926	27117	2.57	6.0E-33	1142919 NT	Homo sapiens similar to RAD23 (S. cerevisiae) homolog B (H. sapiens) (LOC63277), mRNA	
7750	17600	27822	1.24	6.0E-33	675569 NT	Mus musculus SRY-box containing gene 6 (Sox6), mRNA	
7750	17600	27823	1.24	6.0E-33	675569 NT	Mus musculus SRY-box containing gene 6 (Sox6), mRNA	
1739	11640		1.78	5.0E-33 BF 3735 5.1	EST_HUMAN	QY1-F 0169-100700-271-a02 F 0169 Homo sapiens cDNA	
1838	11735		1.18	5.0E-33	11141884 NT	Homo sapiens solute carrier family 5 (choline transporter), member 7 (SLC5A7), mRNA	
1856	11752	21626	1.37	5.0E-33	4507208 NT	Homo sapiens spermidine synthase (SRM) mRNA	
1856	11752	21627	1.37	5.0E-33	4507208 NT	Homo sapiens spermidine synthase (SRM) mRNA	

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Single Exon Probes Expressed in Heart

Probe SEQ ID NO:	Exon SEQ ID NO:	ORF SEQ ID NO:	Expression Signal	Most Similar (Top) Hit BLAST E Value	Top Hit Accession No.	Top Hit Database Source	Top Hit Descriptor
2226	12111			1.3	5.0E-33 AL_163285.2	NT	Homo sapiens chromosome 21 segment HS21C085
9082	18858			1.62	5.0E-33 11433063	NT	Homo sapiens ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome) (UBE3A), mRNA
1112	11027			3.28	4.0E-33 AL_163207.2	NT	Homo sapiens chromosome 21 segment HS21C007
2078	11988	21861		1.77	4.0E-33 4755987	NT	Homo sapiens RAB1, member RAS oncogene family (RAB1) mRNA
2370	12250			7.39	4.0E-33 AA626621.1	EST_HUMAN	ab51b11.1 Stratagene lung carcinoma 93/218 Homo sapiens cDNA clone IMAGE:8443175' similar to contains Alu repetitive element
2498	12373	22265		3.77	4.0E-33 AL_163210.2	NT	Homo sapiens chromosome 21 segment HS21C010
4382	14278	24057		1.63	4.0E-33 AW293349.1	EST_HUMAN	UI-H-BI2-ah1-c-03-U1_s1 NCI_CGAP_Sub4 Homo sapiens cDNA clone IMAGE:2727149'3'
5318	15239	25043		20.94	4.0E-33 AA053053.1	EST_HUMAN	Z71a@3.r1 Stratagene colon (#03/204) Homo sapiens cDNA clone IMAGE:510038 5' similar to gb:X12871_ma1 HETEROGENEOUS NUCLEAR RIBONUCLEOPROTEIN A1 (HUMAN);
9007	18810	29102		1.72	4.0E-33 11425635	NT	Homo sapiens killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1 (KIR3DL1), mRNA
1073	10989			4.46	3.0E-33 BE350127.1	EST_HUMAN	ht09g01.x1 NCI_CGAP_Kid13 Homo sapiens cDNA clone IMAGE:3146256 3' similar to contains MER29,b3
1074	10989			3.4	3.0E-33 BE350127.1	EST_HUMAN	MER29 repetitive element;
2400	12721			1.48	3.0E-33 AV647851.1	EST_HUMAN	ht09g01.x1 NCI_CGAP_Kid13 Homo sapiens cDNA clone IMAGE:3146256 3' similar to contains MER29,b3
99	10003			2.33	2.0E-33 AI160189.1	EST_HUMAN	AV647851 GLC Homo sapiens cDNA clone IMAGE:1705204 3' similar to qb67g03.x1 Scares_fetal_heart_NbHH19W Homo sapiens cDNA clone IMAGE:1705204 3' similar to
4322	14219			4.27	2.0E-33 BE159039.1	EST_HUMAN	contains OFR,r1 ORF repetitive element;
4911	14790	24565		5.1	2.0E-33 AA626683.1	EST_HUMAN	MR0-HT0405-160300-202-d08 HT0405 Homo sapiens cDNA clone IMAGE:844338 5' similar to ab51g11.1 Stratagene lung carcinoma 93/218 Homo sapiens cDNA clone IMAGE:844338 5' similar to
5026	14899	24668		1.73	2.0E-33 11421332	NT	gb:X00734_cds1 TUBULIN BETA-5 CHAIN (HUMAN);
5026	14899	24669		1.73	2.0E-33 11421332	NT	Homo sapiens hypothetical protein SIRP-b2 (SIRP-b2), mRNA
5869	15775	25894		1.61	2.0E-33 AI277492.1	EST_HUMAN	qj96dd1.x1 Scares_NhhMPu_S1 Homo sapiens cDNA clone IMAGE:1880161 3'
7261	17138			1.8	2.0E-33 A1052256.1	EST_HUMAN	oz21d03.x1 Scares_fetal_liver_spleen_1NFLS-S1 Homo sapiens cDNA clone IMAGE:1675973 3' similar to
8	9994			1.44	1.0E-33 AF003528.1	NT	gb:M29536 TRANSLATIONAL INITIATION FACTOR 2 BETA SUBUNIT (HUMAN);
8637	18502	28778		2.02	1.0E-33 AW996618.1	EST_HUMAN	Hom sapiens X-linked anhidrotic ectodermal dysplasia protein (EDA), exon 2 and flanking repeat regions
8921	18729	28024		5.33	1.0E-33 U60822.1	NT	QV3-BN0047-230200-102-b03 BN0047 Homo sapiens cDNA Human dystrophin (DMD) gene, exons 7, 8 and 9, and partial cds
9553	19151			1.92	1.0E-33 A927191.1	EST_HUMAN	w08806.x1 NCI_CGAP_Kid11 Homo sapiens cDNA clone IMAGE:2462410 3'